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(54) Title: VIRAL VECTOR WITH BOVINE VIRAL DIARRHEA VIRUS (BVDV) ANTIGENS			
(57) Abstract <p>This invention relates to the field of Bovine Viral Diarrhea Virus (BVDV), and vaccines for the treatment thereof. This invention describes the preparation of live, attenuated Bovine Herpesvirus type 1 (BHV-1) as a virus, vaccine and vector for expression of BVDV antigens. A BVDV cDNA clone containing sequences corresponding to glycoprotein gp53 is inserted into an inactivated BHV-1 virus.</p>			

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VIRAL VECTOR WITH BOVINE VIRAL DIARRHEA VIRUS (BVDV) ANTIGENS**BACKGROUND OF THE INVENTION**Field of the Invention

5 This invention relates to the field of Bovine Viral Diarrhea Virus (BVDV), and vaccines for the treatment thereof.

Information Disclosure

van Zijl, M. *et al.* Live Attenuated Pseudorabies Virus Expressing Envelope Glycoprotein E1 of Hog Cholera Virus Protects Swine against both Pseudorabies and Hog Cholera, *Journal of Virology*, Vol. 65, No. 5, pp. 2761-2765 (1991). U.S. patent 4,703,011, Kit, M., and Kit, S., Thymidine Kinase Deletion Mutants of Bovine Herpesvirus-1, issued 27 October 1987. U.S. patent 4,824,667, Kit, M., and Kit, S. Thymidine Kinase Deletion Mutants of Bovine Herpesvirus-1, Vaccines Against Infectious Bovine Rhinotracheitis Containing Same and Methods for the Production and Use of Same, issued 25 April, 1989. Collett, M.S., *et al.*, Proteins Encoded by Bovine Viral Diarrhea Virus: The Genomic Organization of a Pestivirus, *Virology*, Vol. 165 pp. 200-208 (1988). Collett, M.S., *et al.*, Molecular Cloning and Nucleotide Sequence of the Pestivirus Bovine Viral Diarrhea Virus, *Virology*, Vol. 165 pp. 191-199 (1988).

20 Background

Bovine viral diarrhea virus (BVDV) is a Pestivirus belonging to the family of the Flaviviridae. It causes a number of different conditions in sheep, goats, and especially cattle. The symptoms depend upon the age, physiological and virological state of the animal. In young susceptible calves and young adults it causes a disease which is characterized by high morbidity and low mortality. The symptoms can include fever, depression, occulo-nasal discharges, diarrhea and occasionally oral ulcerations. Apart from these primary effects the virus also causes immunosuppression. Although primary BVDV infections are normally relatively mild, the virus may potentiate or enhance the pathogenicity of other co-infecting microorganisms.

30 In older or susceptible animals, BVDV causes similar symptoms to those described above for younger susceptible calves. In addition, in pregnant animals the virus has the ability to cross the placenta and infect the fetus. The outcome of this infection depends upon the age of the fetus and whether it is at a stage where its immune system is fully competent. The possible outcome of infections include fetal

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reabsorption, abortion, mummification, congenital defects, birth defects, calves born which are persistently infected with BVDV and completely normal calves. Calves born which are persistently infected with BVDV, represent the most important segment of this BVDV pathogenesis complex. Persistently infected animals shed
5 large amounts of virus into their environment which can infect susceptible animals. Furthermore, even though persistently infected animals are immunotolerant to the virus which infected them in utero, they do develop disease when infected with other closely related BVDV biotypes. These infections are characterized by low morbidity (because relatively speaking there will not be many pregnant animals infected at the
10 right time during pregnancy to produce BVDV persistently infected normal calves), but high mortality. This disease syndrome is known as mucosal disease and often manifests itself as a peracute condition with calves dying of a profuse watery diarrhea which contains large amounts of fresh blood.

The importance of this virus and its widespread presence in the cattle
15 population has led to the development of many vaccines in the attempt to try to prevent BVDV infection. These vaccines have been built on the traditional concepts of inactivation or attenuation but, because of the behavior of BVDV, they have many significant drawbacks.

It is generally accepted that inactivated vaccine preparations are not as
20 effective as attenuated live vaccines. Inactivated antigen from inactivated vaccine preparation undergoes exogenous processing. After injection into the animal the antigen becomes part of the animal's soluble protein milieu. The antigen enters antigen presenting cells through pinocytotic mechanisms and this usually produces antibodies. Unfortunately, because antibodies cannot gain entry into cells, they
25 normally only interrupt viral life cycles when mature virus is released from the cell. On the other hand, antigen from live virus which replicates inside cells, undergoes endogenous processing and this mechanism produces the preferred cell mediated immune responses. Cell mediated immune responses can recognize cells infected with viruses and have the potential of interrupting the virus life cycle at a much
30 earlier stage. Cell mediated responses are thus thought to be extremely important in the immunological defense to many viral infections.

Because of the cell mediated response, attenuated live products such as vaccines should induce good cell mediated responses. With BVDV, attenuation of the virus to produce the live vaccine does not always prevent that vaccine virus from
35 causing the immunosuppression normally associated with field isolates. Roth J.A.

and Kaeberle M.L., Suppression of Neutrophil and Lymphocyte Function Induced by a Vaccinal Strain of Bovine Viral Diarrhea Virus With or Without the Administration of ACTH, *American Journal of Veterinary Research*, Vol. 44 pp. 2366-2372 (1983). The failure of the vaccine to stop the immunosuppression response

5 creates a serious drawback to the vaccine. An animal owner may be vaccinating animals to protect against a disease but because of the properties of the vaccine the owner provides an opportunity for other diseases to afflict the animals. This forces the owner to use inactivated BVDV vaccines, which because of the way in which the immune system operates, are not particularly effective.

10 In summary, inactivated vaccines are safe but not particularly effective while the attenuated live vaccines are more effective but under certain conditions may not be very safe.

This invention combines the effectiveness of the attenuated live vaccines with the safety of the inactivated vaccines. Bovine herpesvirus type 1(BHV-1) is another
15 major pathogen of cattle which produces respiratory disease. Thus, in common with BVDV, BHV-1 also replicates at a mucosal surface. We take the gene which codes for gp53, a major glycoprotein of the BVDV virus and against which the host produces substantial immune responses, and express it in bovine herpes virus -1 (BHV-1), this recombinant virus (BHV/BVDVgp53) is used as a vaccine against
20 BVDV. Donis, R.O. and Dubovi, E.J., Glycoproteins of Bovine Viral Diarrhoea-Mucosal Disease Virus in Infected Bovine Cells, *Journal of General Virology*, Vol. 68, pp. 1607-1616 (1987) and Magar, R., et al., Bovine Viral Diarrhea Virus Proteins: Heterogeneity of Cytopathogenic and Noncytopathogenic Strains and Evidence of 53K Glycoprotein Neutralization Epitope, *Veterinary Microbiology*, Vol. 16, pp. 303-
25 314. Cited references are incorporated herein by reference.

SUMMARY OF THE INVENTION.

A replicating nonpathogenic virus, for preventing disease caused by Bovine Viral Diarrhea Virus (BVDV), where said replicating nonpathogenic virus comprises: a gene or gene combination taken from a BVDV virus, and said replicating
30 nonpathogenic virus functionally expresses said gene or gene combination. Embodiments of this invention include the following: A virus where said replicating nonpathogenic virus is attenuated, is selected from attenuated Bovine Herpes Virus type 1 (BHV-1), attenuated adenoviruses, attenuated bovine mammillitis virus, attenuated bovine papillomavirus, or attenuated pseudorabies virus. A virus where
35 said replicating nonpathogenic virus is attenuated and contains and expresses any

combination of the following genes: the genes that code for gp48, gp25, p14 capsid protein, p20 N-terminal protease and p125/p80 protein. A virus where the attenuation is created by making the thymidine kinase (tk) gene nonfunctional.

- A virus where a signal peptide is inserted preceeding the gene or gene combination that codes for gp53 in said Bovine Herpes Virus type 1 (BHV-1). A virus where said gene that codes for gp53 is inserted into the inactivated thymidine kinase (tk) gene site. A virus where the functionally expressing gene or gene combination, used to create the virus, comprises a recombined plasmid with intact viral DNA, said plasmid comprising: a) a BHV-1 genomic DNA fragment containing the thymidine kinase (tk) gene and having a deletion to the thymidine kinase (tk) gene, b) a promoter/polyadenylation signal inserted in the thymidine kinase (tk) gene deletion, c) a signal peptide gene sequence preceding a gp53 gene or gene combination all of which is inserted between the promoter and the polyadenylation signal. A virus where said plasmid is made from a plasmid having the characteristics of plasmid pHAS4. A virus where said signal peptide gene sequence is taken from any well characterized signal peptide sequences such as any of the thirty-nine examples of well characterized signal peptide sequences found in Perlman, D., et al., *J. Mol. Biol.* Vol. 167 pp. 391-409 (1983), incorporated by reference. A virus where said signal peptide gene sequence is taken from Psuedorabies Virus gIII gene (PRV) and/or Bovine Growth Hormone (BGH).

- A virus where a plasmid is selected from the following plasmids, a) pBHVtkex-1::BGH/p53; b) pBHVtkex-1::gIII/p53; c) pBHVtkex-3::BGH/p53; or d) pBHVtkex-3::gIII/p53. A virus that produces the product of a functionally expressing gene or gene combination is selected from one of the following viruses, T11-3, T11-6, or T11-8. A virus where the functionally expressing gene or gene combination, used to create the virus, comprises a recombined plasmid with intact viral DNA, said plasmid comprising: a) a BHV-1 genomic DNA fragment containing the thymidine kinase (tk) gene and having a deletion to the thymidine kinase (tk) gene, b) a promoter/polyadenylation signal inserted in the thymidine kinase (tk) gene deletion, c) a gp53 gene or gene combination inserted between the promoter and the polyadenylation signal. A virus where the plasmid is pBHVtkex-3::p53. A virus selected from one of the following viruses, T2-3#3 or T2-2#5. A vaccine for preventing disease caused by Bovine Viral Diarrhea Virus (BDVD) comprising a pharmaceutically effective amount of the viruses described herein and a carrier.

- A vaccine as described above for preventing disease caused by Bovine Viral

Diarrhea Virus (BDVD) comprising a pharmaceutically effective amount of a virus described above and a carrier, said carrier comprising any physiological buffered medium, i.e. about pH 7.0 to 7.4 containing from about 2.5 to 15% serum which does not contain antibodies to BHV.

5 A method of immunizing an animal against infectious disease caused by Bovine Viral Diarrhea Virus (BDVD) comprising administering to an animal a pharmaceutically effective amount of a virus or vaccine described herein.

 A process of preparing a virus described herein comprising: a) isolation of a functionally expressing gene or gene combination that causes BVDV, b) inserting
10 said gene or gene combination into a replicating nonpathogenic virus, c) selecting a live-virus that functionally expresses the product of said gene or gene combination.

 A method of preparing a virus described herein where the functionally expressing gene or gene combination, used to create the virus, is produced by a process comprising the recombination of a plasmid with intact viral DNA, said
15 plasmid comprising: a) a BHV-1 genomic DNA fragment containing the thymidine kinase (tk) gene and having a deletion to the thymidine kinase (tk) gene, b) inserting into the thymidine kinase (tk) gene deletion of said plasmid a promoter/polyadenylation signal, c) inserting a gp53 gene or gene combination between the promoter and the polyadenylation signal, d) transfecting cells with said
20 plasmid to produce a recombinant virus containing said functional gene or gene combination inserted into a live virus that does not cause immunosuppression in the usual host and expressing said functional gene or gene combination.

 A method of preparing a virus described herein where the functionally expressing gene or gene combination, used to create the virus, is produced by a
25 process comprising the recombination of a plasmid with intact viral DNA, said plasmid comprising: a) a BHV-1 genomic DNA fragment containing the thymidine kinase (tk) gene and having a deletion to the thymidine kinase (tk) gene, b) inserting into the thymidine kinase (tk) gene deletion of said plasmid a promoter/polyadenylation signal, c) inserting a gp53 gene or gene combination
30 preceded by a signal peptide gene sequence between the promoter and the polyadenylation signal, d) transfecting cells with said plasmid to produce a recombinant virus containing said functional gene or gene combination inserted into a live virus that does not cause immunosuppression in the usual host and expressing said functional gene or gene combination.

BRIEF DESCRIPTION OF THE DRAWINGS.

- Figure 1. Construction of the shuttle vectors for inserting foreign genes into BHV-1.
 Figure 2. Strategy for appending signal peptide sequences to the BVDV gp53 gene.
 Figure 3. Maps of the five shuttle plasmids for inserting gp53 into BHV-1
- 5 a. EXAMPLE 1. pBHVtkex-3::p53.
 - b. EXAMPLE 2: pBHVtkex-1::BGH/p53
 - c. EXAMPLE 3: pBHVtkex-1::gIII/p53
 - d. EXAMPLE 4: pBHVtkex-3::BGH/p53
 - e. EXAMPLE 5: pBHVtkex-3::gIII/p53
- 10 Figure 4. Predicted transcript maps of the BHV-1/gp53 recombinant viruses.
 Figure 5. Northern blots showing transcription of gp53 messenger RNAs in the
 BHV-1 recombinants.
 Figure 6. Immunoprecipitations showing expression of gp53 protein in the BHV-1
 recombinants.

15 DESCRIPTION OF THE PREFERRED EMBODIMENTS.

All of the terms used below will be readily understood by one skilled in the
 art. In many places the name of the manufacturer of equipment or reagents are
 provided in parenthesis after the equipment or reagent is named. Commonly used
 terms, reagents and buffers such as "plasmids," "Klenow Fragments," "religating
 20 blunt ends," "Tris," chelating buffers such as EDTA and EGTA, and commonly used
 chromatography columns are referred to without further explanation.

In the descriptions of the construction of the compounds used in this
 invention, standard molecular biological techniques were used and are briefly named
 or described here. Detailed explanations of these techniques can be found in
 25 standard laboratory manuals such as "Molecular Cloning: a Laboratory Manual"
 (1989), Sambrook, *et. al.*, Cold Spring Harbor Press, Cold Spring Harbor, New York,
 or "Current Protocols in Molecular Biology" (1991), Ausubel, F. M., *et. al.*, eds., Wiley
 Interscience, New York.

This invention combines the effectiveness of the attenuated live vaccines with
 30 the safety of the inactivated vaccines. We take the gene which codes for gp53, a
 major glycoprotein of the BVDV virus and against which the host produces
 substantial immune responses, and express it in bovine herpesvirus -1 (BHV), this
 recombinant virus (BHV/BVDVgp53) is used as a vaccine against BVDV.

Bovine herpesvirus (BHV) is another major pathogen of cattle which produces
 35 respiratory disease. Thus, in common with BVDV, BHV also replicates at a mucosal

surface. With BVDV, replication is mainly at the gut mucosal interface with less replication at the respiratory interface. With BHV it is the respiratory interface which dominates. The common mucosal immune system ensures that immune responses produced at one surface will be effective at other surfaces. Thus the recombination virus of this invention, BHV/BVDVgp53, will, when administered to cows, sheep or goats, preferably via the intranasal route, replicate in the respiratory mucosae and produce an immune response.

Prior to the expression of the BVDVgp53 gene in BHV, the thymidine kinase gene was deleted from the BHV virus using a process known to attenuate the virus. The BHV, a live attenuated virus, will replicate and produce a cell mediated response. As part of that replicative process, the BVDV gp53 gene will be expressed and, because the virus is inside the cell, the correct processing for a cell mediated response to the BVDV gp53 part of the recombinant virus will also occur. Most importantly, this response will occur without the possible side effects of immunosuppression, as only part of the BVDV virus is present. Thus, the invention combines the efficacy of an attenuated live virus vaccine for BVDV, with the safety of an inactivated preparation.

The examples in the procedures section are provided for illustrative purposes and are in no way intended to limit the scope of the present invention. All media and buffer solutions were made up in glass distilled water unless otherwise indicated.

Compositions and Administrations - A pharmaceutically effective amount of the vaccine of the present invention can be employed along with a pharmaceutically acceptable carrier or diluent as a vaccine against BHV-1 and BVDV in animals, such as bovine, sheep and goats.

Examples of pharmaceutically acceptable carriers or diluents useful in the present invention include any physiological buffered medium, i.e., about pH 7.0 to 7.4, containing from about 2.5 to 15% serum which does not contain antibodies to BHV, i.e., is seronegative for BHV. Serum which does not contain gamma globulin is preferred to serum which contains gamma globulin. Examples of serum to be employed in the present invention include fetal calf serum, lamb serum, horse serum, swine serum, and goat serum. Serum protein such as porcine albumin or bovine serum albumin (hereinafter "BSA") in an amount of from about 0.5 to 3.0% can be employed as a substitute for the serum. However, it is desirable to avoid the use of foreign proteins in the carrier or diluent which will induce allergic responses

in the animal being vaccinated.

The virus may be diluted in any of the conventional stabilizing solutions containing phosphate buffer, glutamate, casitone, and sucrose or sorbose, or containing phosphate buffer, lactose, dextran and glutamate.

5 It is preferred that the vaccine viruses of the present invention be stored at a titer of at least 10^5 to 10^6 PFU/ml at -70°C to -90°C or in a lyophilized state at 2°C to 7°C . The lyophilized virus may be reconstituted for use with sterile distilled water or using an aqueous diluent containing preservatives such as gentamicin and amphotericin B or penicillin and streptomycin.

10 The useful dosage to be administered will vary depending upon the age, weight and species of the animal vaccinated and the mode of administration. A suitable dosage can be, for example, about $10^{4.5}$ to 10^7 PFU/animal, preferably about $10^{4.5}$ to $10^{5.5}$ PFU.

The vaccines of the present invention can be administered intranasally,
15 intravaginally or intramuscularly. Intranasally is the preferred mode of administration.

Utility of the Invention - This invention is intended to provide the user with an effective vaccine for prevention of BVDV caused disease, where the vaccine can be safely and efficaciously administered intramuscularly, intranasally, or
20 intravaginally. Intranasally may be the preferred route of administration.

The vaccines of this invention are created with the intention of treating disease, preferably through prevention. By prevent or prevention applicant means to keep the host from developing symptoms of the disease or to mitigate the effects of the disease, that is to avert the typical diseased state. Prevention implies decisive
25 action to stop, impede or delay the onset of disease. Prevention can include the following concepts: to hinder, frustrate, to obstruct; to intercept, possibly prohibit, impede or preclude. Preclude would suggest the onset of the disease state either does not occur or the disease pathogen is largely ineffectual in causing the disease state. Prevent or prevention can indicate the disease state is forestalled, meaning
30 that anticipatory action to prevent or hinder the disease has occurred but the conditions creating the disease have not been eliminated.

The usefulness of this invention will be illustrated by the ability of the vaccine to provide effective protection against the spread of BVDV disease in its various manifestations. Because the vaccine uses gp53, a major glycoprotein of
35 BVDV, and one against which the host produces a substantial immune response, the

vaccine will confer substantial benefits upon the treated potential host. Another object of the invention is to provide a BVDV vaccine which can be administered safely to calves and to pregnant cows in all stages of pregnancy.

Measures of Activity - The vaccine uses gp53, a major glycoprotein of BVDV, and one against which the host should produce a substantial immune response. Others have shown that gp53 is highly immunogenic. Donis, R.O. and Dubovi, E.J., Glycoproteins of Bovine Viral Diarrhoea-Mucosal Disease Virus in Infected Bovine Cells, *Journal of General Virology*, Vol. 68, pp. 1607-1616 (1987). It is well known that agents that produce substantial immune responses can make effective vaccines. Magar, R., *et al.*, Bovine Viral Diarrhea Virus Proteins: Heterogeneity of Cytopathogenic and Noncytopathogenic Strains and Evidence of 53K Glycoprotein Neutralization Epitope, *Veterinary Microbiology*, Vol. 16, pp. 303-314. The vaccines of this invention contain genes that express large quantities of gp53, this is shown in figure 5. Because of the expression of large quantities of gp53 the vaccines of this invention will confer substantial benefits upon the treated potential host.

Preferred Compounds - Any BHV-1 virus attenuated with a tk deletion and carrying the gp53 gene, the gp53 gene being preceded by a signal peptide, that expresses abundant amounts of gp53, should be a preferred suitable vaccine candidate. It appears the signal peptide sequence may be taken from any suitable source. We chose to examine two different signal peptides to ensure the best localization of the gp53 protein *in vivo*. We chose two candidates we call "T11-6", embodied in Example 2, and "T11-3", embodied in Example 3 for vaccine trials. The former virus was deposited to the ATCC under the designation UC VR-58. The latter, "T11-3" plasmid was also deposited. The virus we labeled "T11-8" might contain truncated forms of the tk transcript and this might suggest, but does not necessarily mean, that it would be less attractive as a vaccine candidate. A large number of existing cell lines are persistently infected with non-cytopathic BVDV from passage in media containing fetal bovine serum taken from infected calves. For this invention, it is imperative that viruses used as live, attenuated vaccines are free of contaminating BVDV.

Preparation of the Compounds

Construction of expression shuttle vectors for gene insertion into Bovine herpesvirus type-1 (BHV-1).

We constructed two shuttle vectors to allow insertion of foreign genes into BHV-1. Although this invention shows the utility of BHV-1 as a vector for BVDV

genes many other viruses could fill the same role. Other examples from cattle, sheep and goats would include cow, goat and sheep pox viruses, adenoviruses, bovine mammillitis virus, bovine papillomavirus, and pseudorabies virus. A non-pathogenic virus refers to any virus which has the ability to replicate in one of its host species but does not produce any signs of disease in that species. Such non-pathogenic viruses might arise from pathogenic parent viruses by natural mutation, might be mutagenized by, for instance, chemicals or light to produce a non-pathogenic virus, or could be rendered non-pathogenic through the use of recombinant DNA technologies. See, 1) Mapping Neutralization Domains of Viruses, E. Wimmer, E.A. Emini, and D.C. Diamond and 2) Immunogenicity of Vaccine Products and Neutralizing Antibodies, E. Norrby. Both articles are in Edited by Notkins and Oldstone Published by Springer-Verlag New York Inc. 1986.

Since we intended to attenuate BHV-1 by inactivating the viral thymidine kinase (tk) (M. Kit, et al., US Patent 4,703,011, (1983)), we decided to use the BHV-1 tk gene for the site of insertion. This approach not only insured the complete inactivation of the viral tk, but also allowed us to select recombinant, tk-negative virus by established methods. M. F. Shih, et al., *Proc Natl Acad Sci USA*, 81:5867-5870 (1984). Other methods to attenuate BHV-1, such as deletion of other non-essential genes would also be applicable to this particular invention. We started with plasmid pHAS4 which contains a 2.7kb SalI subfragment of the BHV-1 HindIII-A fragment cloned into plasmid pUC18. E. Petrovskis, unpublished data. M. Engels, et al., *Virus Res*, 6:57-73 (1986); J. E. Mayfield, et al., *J Virol*, 47:259-264 (1983); A. L. Meyer, et al., *Biochim Biophys Acta*, 1090:267-9 (1991). As shown in Fig. 1, this SalI fragment contained the entire tk gene, as well as a portion of the upstream gene homologous to the HSV-1 UL24 gene, and a portion of the glycoprotein H gene. L. J. Bello, et al., *Virology*, 189:407-414 (1992); J. G. Jacobson, et al., *J Virol*, 63:1839-1843 (1989); M. Kit, et al., US Patent 4,703,011, (1983); A. L. Meyer, et al., *Biochim Biophys Acta*, 1090:267-9 (1991).

A 424bp deletion was introduced into the tk gene by digesting pHAS4 with BglII and XhoI, filling in the ends with the Klenow Fragment of DNA polymerase I (Klenow) and religating the resulting blunt ended fragments. This manipulation restored the BglII recognition site, but not the XhoI site (Fig. 1). The resulting plasmid was named pHAS4ΔBX. This deletion was chosen because it does not impede on the previously identified transcription initiation sites for the UL24 homolog which overlaps the 5' end of the tk gene. L. J. Bello, et al., *Virology*,

189:407-414 (1992); J. G. Jacobson, et al., *J Virol*, 63:1839-1843 (1989). Numerous other deletions within the BHV-1 tk gene would be possible. To facilitate later cloning manipulations, we eliminated the HindIII site in the pUC18 vector by digesting pHAS4ΔBX with HindIII, filling in the cohesive ends with Klenow, and
 5 religating the blunt ends.

We obtained a 1775bp cassette containing the Human cytomegalovirus (CMV) major immediate early promoter and the bovine growth hormone polyadenylation sequence. R. J. Brideau, et al., *J Gen Virol*, 74:471-477 (1993). These gene expression signals are commonly used for high levels of expression of foreign genes
 10 in a number of different systems, but other promoter/polyadenylation signal pairs could also be used in this context. The cassette, in vector p3CL-DHFR, is bounded by unique EcoRI and BglII sites and contains, between the promoter and the polyadenylation signal, unique HindIII and Sall restriction sites for cloning of foreign genes. The p3CL-DHFR vector was digested with EcoRI, then filled in and
 15 ligated to a BamHI linker (New England Biolabs, Beverly, Massachusetts). This manipulation regenerated the EcoRI site. The construct was then digested with BamHI and BglII and the released cassette was ligated into the BglII site of pHAS4ΔBX (Fig. 1). The ligations were transformed into *E. coli* strain DH5α. We isolated recombinant plasmids that contained the p3CL insert in both orientations
 20 relative to the BHV-1 tk gene by mapping of asymmetric restriction sites. These two constructs, designated pHAS4ΔBXex-1 and pHAS4ΔBXex-3 (Fig. 1), contained then, a strong promoter and polyadenylation signal bounded by the BHV-1 tk gene and flanking regions to allow homologous recombination into the BHV-1 genome.

Figure 1. Construction of shuttle vectors for inserting foreign genes into
 25 BHV-1. PHAS4 is a 2.7kb subfragment from the BHV-1 HindIII-A fragment. The BglII/XhoI subfragment to be deleted is shown. The deletion derivative of pHAS4 is pHAS4ΔBX. The deleted thymidine kinase (tk) gene is shown as a dark stippled box. The cassette containing the promoter and polyadenylation signal is shown just below pHAS4ΔBX. The CMV immediate early promoter is shown as a light stippled
 30 box, and the Bovine Growth Hormone (BGH) polyadenylation signal is shown as a striped box. Finally, the inserts of the two expression shuttle plasmids, pHAS4ΔBXex1 and pHAS4ΔBXex3 are shown.

Addition of Signal Peptide Sequences to BVDV gp53 gene.

A cDNA containing the BVDV gp53 gene from strain 2724, a noncytopathic
 35 strain, has been previously described. Kennedy, M. et al, abstracts of the American

College of Veterinary Microbiologists, 1992 workshop. Since the BVDV RNA genome is normally translated as one long polypeptide and then post-translationally modified into the various viral proteins, the gp53 portion of the BVDV genome does not contain the usual signal peptide required for translocation of the protein to the cell membrane, where the protein is normally expressed. Nonetheless, the cDNA was successfully expressed in both cell-free systems and baculovirus, and the protein appeared to be translocated, glycosylated and anchored in both systems, despite the lack of a conventional signal peptide. We decided, however, to evaluate expression of gp53 in BHV-1 both with and without various signal peptides.

10 In order to attach nucleotide sequences encoding signal peptides to the gp53 gene, we introduced a BamHI site into 5' end of the p53 gene by site directed mutagenesis, as follows: The p53 gene was blunt-end ligated into the filled-in BamHI site of plasmid pSP72 (Promega Corp., Madison, Wisconsin), thus removing all BamHI sites from the resulting plasmid. We introduced a single base change, a C to a G, 11 bases in from the initiation codon used by the cDNA, using a synthetic oligonucleotide and the "Double Take" site directed mutagenesis kit (Stratagene, La Jolla CA) according to the manufacturer's instructions. This base change introduced a unique BamHI site into the gene without altering the gp53 amino acid sequence (Fig. 2 section B). The base change was verified by nucleotide sequencing, and the resulting plasmid was called pP53mut. We inserted, into pP53mut sequences, encoding signal peptides from the PRV gIII gene (A. K. Robbins, et al., *J Virol*, 58:339-347 (1986)) and from Bovine growth hormone. R. P. Woychik, et al., *Nucl Acids Res*, 10:7197-7210 (1982). (Figure 2 section A) Complementary oligonucleotides encoding the two signal peptides were synthesized such that annealed oligos had SalI cohesive ends 5' and BamHI cohesive ends 3' (Fig 2 section A). These signal peptide cassettes were ligated into pP53mut digested with BamHI and SalI, and transformed into DH5 α . We confirmed the correct insertion of the signal peptide cassettes by nucleotide sequencing.

30 Complementary oligonucleotides encoding any well characterized signal peptide can be used in this invention. Thirty-nine examples of well characterized signal peptide sequences found in Perlman, D., et al., *J. Mol. Biol.* Vol. 167 pp. 391-409 (1983). Incorporated by reference. These and any other well characterized signal peptides should be suitable for use as embodiments of this invention.

Figure 2. Strategy for appending signal peptide sequences to the BVDV gp53 gene. Section A: Synthetic oligonucleotides corresponding to the signal

peptide sequences of Bovine Growth Hormone (BGH), and Pseudorabies virus gIII (PRV gIII). Complementary oligonucleotides were synthesized such that the annealed pairs had SalI sites on the 5' ends and BamHI sites on the 3' ends. The deduced amino acid sequences of the signal peptides are also shown. In each case
5 the predicted cleavage sites for the signal peptides are just after the alanine (A), three amino acids from the ends. Codons for two amino acid residues (F,P in BGH; P,S in gIII) from the original native proteins were left on the signal peptide sequences to ensure correct cleavage.

Section B: Site directed Mutagenesis of the cDNA encoding the BVDV gp53
10 gene. The first 60 nucleotides of the gp53 cDNA and the corresponding amino acid sequence are shown. A single base pair, shown by the arrow, was changed to create a BamHI restriction site in the sequence, shown in the box. This change does not change the amino acid sequence. The cDNA was then digested with BamHI as shown, allowing in frame ligation to either of the signal peptide sequences shown in
15 section A.

Other expression gene fragments in addition to gp53.

Expression of other BVDV gene or gene combinations in a live virus vector are also embodiments of this invention. This would include any and all BVDV proteins to which a vaccinated animal could elicit an immune response. Examples
20 include, but are not limited to, the other two BVDV surface glycoproteins, gp48 and gp25 (Collett, M.S., et al., *Virology* 165:200-208 (1988)), the p14 capsid protein (Thiel, H.J., et al., *J. Virol.* 65:4705-4712 (1991)), and the p20 N-terminal protease. Wiskerchen, M., et al., *J. Virol.* 65:4508-4514 (1991). This group of proteins, along with the gp53 gene, can be expressed together from a single cDNA molecule, the
25 expressed polyprotein will process itself correctly into the separate proteins. Another BVDV protein candidate to express in a vaccine is the nonstructural p125/p80 protein (Deregt, D., et al., *Can. J. Microbiol.* 37:815-122 (1991)), which elicits a significant antibody response in infected cows.

Insertion of the BVDV gp53 gene into the BHV-1 expression vectors.

30 The p53 gene, either with or without added signal peptide sequences, was ligated into the HindIII insertion sites of pHAS4ΔBXex-1 and pHAS4ΔBXex-3 by filling in all the respective cohesive ends of vectors and inserts followed by blunt end ligation. The ligations were transformed in *E. coli* strain DH5α. We wanted to eventually evaluate the expression of gp53 in BHV-1 in various orientations and
35 with at least two different signal peptides to ensure that we achieved the most

efficient expression. The transformed colonies were screened by colony hybridization using as a probe the p53 insert labelled with Digoxigenin-dUTP. The "Genius" DNA hybridization system (Boeringer Mannheim Biochemicals (BMB), Indianapolis, IN) was used for this and all other DNA hybridizations described in the characterization of this invention. Positive recombinants were then screened by restriction analysis for those carrying the gp53 gene in the proper orientation relative to the CMV promoter and BgH polyadenylation signal. Five plasmids were isolated, which are schematically depicted in Figs. 3A-E. Their descriptions are as follows.

EXAMPLE 1. pBHVtkex-3::p53: contains the BVDV gp53 gene inserted between the CMV promoter and the BGH polyadenylation signal of pHAS4ΔBXex-3 with no added signal peptide. In this construct the original gp53 gene, PRIOR to site directed mutagenesis, was inserted. See Fig. 3A. This plasmid was then used to construct the virus T2-3#.

EXAMPLE 2. pBHVtkex-1::BGH/p53: contains the mutagenized gp53 gene preceded by the BGH signal peptide sequence inserted into pHAS4ΔBXex-1. See Fig. 3B. This plasmid was used to create the virus T11-6. This virus was deposited.

EXAMPLE 3. pBHVtkex-1::gIII/p53: contains the mutagenized gp53 gene preceded by the PRV gIII signal peptide sequence inserted into pHAS4ΔBXex-1. See Fig. 3C. This plasmid was used to create the virus T11-3. This plasmid was deposited.

EXAMPLE 4. pBHVtkex-3::BGH/p53: contains the mutagenized gp53 gene preceded by the BGH signal peptide sequence inserted into pHAS4ΔBXex-3. See Fig. 3D.

EXAMPLE 5. pBHVtkex-3::gIII/p53: contains the mutagenized gp53 gene preceded by the PRV gIII signal peptide sequence inserted into pHAS4ΔBXex-3. See Fig. 3E. This plasmid was used to create the virus T11-8. This plasmid was deposited.

Figures 3A-E. Complete maps of the five shuttle plasmids for inserting gp53 into BHV-1. The gp53 gene is shown as a solid band, the BHV-1 sequences are shown as dark stippled bands, the CMV promoter region is shown as a light stippled band, and the BGH polyadenylation signal region is shown as a striped band. The plasmid vector, pUC18, is shown as a thin line. In each case the direction of transcription of gp53 relative to the original direction of transcription of BHV-1 tk is shown. The various signal peptide sequences are indicated.

- a. EXAMPLE 1. pBHVtkex-3::p53.
- b. EXAMPLE 2. pBHVtkex-1::BGH/p53
- c. EXAMPLE 3. pBHVtkex-1::gIII/p53
- d. EXAMPLE 4. pBHVtkex-3::BGH/p53
- e. EXAMPLE 5. pBHVtkex-3::gIII/p53

5

These, and all other possible insertions of the BVDV gp53 gene into the BHV-1 tk gene are embodiments of this invention. These plasmids and any plasmids created in this manner are known as "Principal Plasmid Vectors" and are the plasmid vectors used to create the virus vaccines of this invention.

10

Introduction of the gp53 gene into BHV-1 "Iowa".

The five expression shuttle plasmids carrying gp53 were linearized by XbaI and cotransfected into Bovine Turbinate (BT) cells with unit length DNA from BHV-1 strain Iowa (tk positive) by the standard CaPO_4 method (R. L. Graham, et al., *Virology*, 52:456-467 (1973)) as modified by Cai (W. Cai, et al., *J Virol*, 61:714-721 (1987)). The cells were obtained from ATCC. The transfections were then subjected to two rounds of selection either on 143tk⁻ cells (S. K. Mittal, et al., *J Gen Virol*, 70:(1989)), or on Rab (BU) cells (S. Kit, et al., *Virology*, 130:381-389 (1983)) in the presence of 100ug/ml 5-Bromo-2'-Deoxyuridine (BDUR, Sigma Chemical Company, St. Louis, Missouri) to isolate virus no longer expressing tk. This is a standard procedure described previously. M. Kit, et al., US Patent 4,703,011, (1983). Other tk⁻ cell lines permissive for growth of BHV-1 can also be used. After the two rounds of BDUR passage, transfections that still showed cytopathic effect were infected onto BT cells under complete media with 1% low melting agarose to obtain single plaques. Multiple single plaques were picked from each transfection and the viral DNAs were screened for the p53 gene by dot-blot DNA hybridization. Although not all transfections survived the BDUR passages (particularly those on the 143 tk⁻ cells, as these cells are only marginally permissive for BHV-1 viral growth), those that did survive yielded 100% recombinant virus. Four different recombinant viruses were isolated and further characterized:

30

EXAMPLE 1. T2-3#3 and T2-2#5 (two identical, but independently isolated viral clones): BHV-1 "Iowa" into which the insert sequences contained in pBHVtkex-3::p53 recombined. Contains the BVDV gp53 gene with no added signal peptide sequence situated between the CMV promoter and the BGH polyadenylation signal, with transcriptional orientation in the same direction as the BHV-1tk gene.

35

EXAMPLE 2. T11-6 (This virus was submitted to ATCC under the designation UC VR-58): BHV-1 "Iowa" into which the insert sequences contained in pBHVtkex-1::BGH/p53 recombined. Contains the BVDV gp53 gene with the BGH signal peptide sequence situated between the CMV promoter and the BGH polyadenylation signal, with transcriptional orientation in the opposite direction relative to the BHV-1 tk gene.

EXAMPLE 3. T11-3: BHV-1 "Iowa" into which the insert sequences contained in pBHVtkex-1::gIII/p53 recombined. Contains the BVDV gp53 gene with the PRV gIII signal peptide sequence situated between the CMV promoter and the BGH polyadenylation signal, with transcriptional orientation in the opposite direction relative to the BHV-1 tk gene.

EXAMPLE 5. T11-8: BHV-1 "Iowa" into which the insert sequences contained in pBHVtkex-3::gIII/p53 recombined. Contains the BVDV gp53 gene with the PRV gIII signal peptide sequence situated between the CMV promoter and the BGH polyadenylation signal, with transcriptional orientation in the same direction as the BHV-1 tk gene.

A virus was not isolated from cotransfections with "Iowa" DNA and plasmid pBHVtkex3::BGH/p53, **EXAMPLE 4**, but this prophetic virus, could be easily created, it and any other BHV-1 viruses containing the BVDVgp53 gene inserted into thymidine kinase gene are embodiments of this invention. We purified DNA from each of these viruses and checked for the proper insertions in the proper orientations by Southern Hybridization using both the gp53 gene and the CMV promoter/BgH polyadenylation cassette as probes (data not shown). All four of the viruses carried the complete promoter/gene/polyadenylation cassettes in the BHV-1 tk gene, deleted as predicted, based on restriction fragment sizes. As a control, with these transfections, we also transfected the pHAS4ΔBX plasmid with BHV-1 "Iowa" unit length DNA and isolated a tk-negative progeny carrying the 424bp deletion in tk (also verified by Southern Hybridization). This virus is named IowaΔBX. All of these viruses were plaque purified twice by limiting dilution on BT cells.

A large number of existing cell lines are persistently infected with non-cytopathic BVDV from passage in media containing fetal bovine serum taken from infected calves. For this invention, it is imperative that viruses used as live, attenuated vaccines are free of contaminating BVDV. In order to ensure that the BHV-1 viruses carrying the BVDV sequences were not contaminated with non-cytopathic BVD virus, we prepared DNA from each of the viruses (including the

parent strain Iowa and Iowa Δ BX) and subjected the DNA preps to extensive RNase treatment using a cloned RNase (RNase ONE, Promega Corporation, Madison, Wisconsin). Since BVDV has only RNA as its genetic material, this manipulation should eliminate any possible contaminating BVDV sequences from the viral DNA preps. We then transfected these RNased viral DNAs into certified BVD-free MDBK cells (ATCC) and picked virus plaques from the transfections to use in further manipulations.

Transcriptional analysis of the gp53 recombinants.

We prepared RNA from each of the recombinant viruses and the parent BHV-1 strain Iowa and evaluated transcription of gp53 by Northern hybridization. A diagram of the possible message species and the probes used is shown in Fig. 4.

Figure 4. Predicted transcripts of the BHV-1/gp53 recombinant viruses later shown in Figure. 5. The two probes are 1) the gp53 cDNA and 2) the SalI/BglII portion of pHAS4 (shown above the maps). The first map shows the predicted transcripts from viruses T11-3 and T11-6, and the second map shows the predicted transcripts from T11-8. The sites of transcript initiation for tk and UL24 are shown for reference.

All of the gp53 recombinant viruses made a 1.6kb message that hybridized with a 32 P-labelled gp53 probe, the size predicted for transcription initiation at the CMV promoter and termination at the BgH polyadenylation site, Fig. 5, probe 1. The T2-3#3 and T2-2#5 virus are not shown. As additional major bands, T11-3 and T11-6 made an 8.5kb transcript and T11-8 and T2-3#3 made a 5.6kb transcript. These transcripts were unique to the recombinant viruses, and were consistent with messages initiating at the CMV promoter, reading through the BgH polyadenylation signal and terminating at the UL24 or tk/gH polyadenylation signals, respectively.

Hybridization with the upstream and downstream probes confirmed the identity of these longer messages. The p53 probe did not hybridize to Iowa, Ia Δ BX or mock infected RNAs. As a quantitation control we used probe pHAS6, an 867bp salI fragment that maps downstream of the tk open reading frame and is internal to the gH gene. A. L. Meyer, et al., *Biochim Biophys Acta*, 1090:267-9 (1991). All of the viruses made equivalent amounts of the 3.1kb gH message (data not shown). This probe also hybridized to the longer p53 messages in T11-8 and T2-3-3, and to the 4.3kb tk message in Iowa, which is 3' coterminal with the gH transcript. L. J. Bello, et al., *Virology*, 189:407-414 (1992).

To examine the transcription patterns upstream of the gp53 insertions, we

used a probe that consisted of the pHAS4 fragment from the upstream SalI site to the BglII site in the tk gene, the beginning of the deletion in the recombinant viruses (probe 2). All of the viruses made a message of approximately 4.4kb which we deduced to be UL24 (Fig. 5, probe 2). This message, however, was smaller than the 5.2kb UL24 message in BHV-1 strain Cooper described by Bello, et al (L. J. Bello, et al., *Virology*, 189:407-414 (1992)) and comigrated with the tk message in the wild-type strain Iowa. Although we did not evaluate these comigrating messages further by using single stranded probes, we detected a tk transcript of 4.2 kb only in the Iowa DNA with probe pHAS6 and we detected similarly sized transcripts in all the viral RNAs with the upstream probe, even though these other viruses cannot be making a wild-type sized tk transcript. In T11-3 and T11-6, the upstream probe did not detect any truncated forms of tk message and hybridized to only the UL24 message and the the 8.5kb p53 message. In T11-8, on the other hand, the probe hybridized to four additional (minor) bands of approximately 5.0, 3.7, 1.8, and 1.0kb.

Figure 5. Northern blots showing transcription of gp53 messenger RNAs in the BHV-1 recombinant viruses. The first panel shows transcripts hybridizing to probe 1, the pg53 cDNA, and the second panel shows transcripts hybridizing to probe 2, the SalI/BglII subfragment of pHAS4. KEY: M=Mock infected cells, I=BHV-1 "Iowa" infected cells, 3,6,8=T11-3, T11-6 and T11-8 infected cells. RNA size standards, in kilobases (kB) are given to the left of each panel.

Expression of BVDV gp53 protein in BHV-1.

We evaluated expression of gp53 protein in the BHV-1 recombinants by immunoprecipitation (IP). Detailed procedures for IPs can be found in standard references such as "Current Protocols in Molecular Biology", Ausubel, F. M., et. al., eds., Wiley Interscience, New York. BT cells infected with the BHV-1 recombinants were metabolically labelled with ³⁵S-methionine (Amersham, Arlington Heights, Illinois). The viral infected cells were lysed and soluble proteins were reacted with hyperimmune serum from bovine or goat against BVDV. VMRD, Pullman, Washington. Antigen/antibody complexes were precipitated with staph A (Immunoprecipitin, Gibco/BRL, Gaithersburg, Maryland,) or protein A sepharose 4B (Pharmacia, Uppsala, Sweden). Immunoreactive proteins were resolved by SDS-Polyacrylamide gel electrophoresis (SDS-PAGE) and fluorography.

Figure 6 shows that all three of the recombinant viruses carrying the gp53 gene preceded by a signal peptide sequence made significant amounts of the protein.

We did not detect any expression of gp53 from T2-3#3, or T2-2#5 the viruses carrying the gp53 gene, but lacking a signal peptide, even though this virus synthesized considerable amounts of gp53 messenger RNA. The clones t2-3#3 and T2-2#5 are independently isolated clones, which rules out the possibility that one particular virus had a defect that precluded gp53 expression (data not shown). The possibility remains that gp53 is being synthesized from T2-3, but is rapidly degraded, or that our antibody does not detect unprocessed forms of the protein.

Figure 6. Immunoprecipitated proteins showing expression of gp53 in the BHV-1 recombinants. Labelled proteins were precipitated with polyclonal bovine-anti-BVDV serum, this serum also had minor reactivity with BHV-1 antigens. KEY: 3,6,8=T11-3, T11-6, and T11-8 infected cell proteins, IA=BHV-1 "Iowa" infected cell proteins, M=Mock infected cell proteins. MW=approximate protein molecular weight standards, in Kilodaltons.

The gp53 protein bands in T11-3, T11-6 and T11-8 were broad, suggesting that the proteins were processed, and they appeared to be equivalent but not identical in size to the gp53 protein in NADL (data not shown). Removal of the N-linked sugars from the BVDV-NADL and BHV-1 expressed gp53 proteins by digestion with N-glycansase (Genzyme, Cambridge, Massachusetts) did not resolve the size difference in the proteins, but the proportional reduction in size of the proteins suggested that the native and recombinant forms of gp53 were processed similarly. The slight size difference between the recombinant and native proteins could be due to the fact that the gp53 gene in the BHV-1 viruses came from a different BVD strain which could have a gp53 of a slightly different size, or the cDNA gp53 clone might not contain the exact amino acids processed from the BVDV polypeptide into native gp53.

The present invention is not to be limited in scope by the cell lines deposited or the embodiments disclosed herein which are intended as single illustrations of one aspect of the invention and any which are functionally equivalent are within the scope of the invention. Indeed, various modifications of the invention, in addition to those shown and described herein, will become apparent to those skilled in the art from the foregoing description. Such modifications are intended to fall within the scope of the appended claims.

It is also to be understood that all base pair and amino acid residue numbers and sizes given for nucleotides and peptides are approximate and used for the purposes of description.

All documents cited herein are incorporated by reference.

Deposit of Genetic Materials

One skilled in the art should be able to reconstruct all the various embodiments of this invention by utilizing only the written description. However, for the sake of completeness, to ensure enablement, and to provide every opportunity for others to make and use this invention, certain genetic constructs of this invention have been deposited at recognized depositories in accordance with the Budapest Treaty.

A virus was deposited with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland, zip code 20852, USA. That deposit was designated UC VR-58 by the Upjohn Company and given the following number by the depository, ATCC No. VR2436, it corresponds to the virus described herein as "T11-6," also known as "Example 2." This deposit was received by the American Type Culture Collection depository on 28 October 1993.

Several plasmids were deposited with the Agricultural Research Service Culture Collection (NRRL), of the U.S. Department of Agriculture, at 1815 North University Street, Peoria, Illinois, zip code 61604, USA. One plasmid was given the Upjohn designation, pUC 1564, *E. coli* culture UC 15085, referring to pBHVtkex-1::gIII\p53, it corresponds to the plasmid used to create the virus described herein as "T11-3," also known as "Example 3." This plasmid was given the following number by the depository, NRRL B-21350. Another deposit was given the Upjohn designation, pUC 1565, *E. coli* culture UC 15086, referring to pBHVtkex-3::gIII\p53, it corresponds to the plasmid used to create the virus described herein as, "T-11-8," also known as "Example 5." This plasmid was given the following number by the depository, NRRL B-21351. Both of the plasmids were received by the Agricultural Research Service Culture Collection depository on 26 October 1994.

SEQUENCE LISTING

- (1) GENERAL INFORMATION:
- 5 (i) APPLICANT: The Upjohn Company
INVENTORS (For U.S. Purposes only): Wardley, Richard C. and
Haanes, Elizabeth J.
- 10 (ii) TITLE OF INVENTION: A Replicating Nonpathogenic Virus Expressing
Envelope Glycoproteins from Bovine Viral Diarrhea Virus (BVDV)
- (iii) NUMBER OF SEQUENCES: 2
- 15 (iv) CORRESPONDENCE ADDRESS:
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(C) CITY: Kalamazoo
20 (D) STATE: Michigan
(E) COUNTRY: U.S.A
(F) ZIP: 49001-0199
- 25 (v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- 30 (vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:
- 35 (viii) ATTORNEY/AGENT INFORMATION:
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(C) TELEX: 224 401 UPJOHN
- 45 (2) INFORMATION FOR SEQ ID NO:1:
- (i) SEQUENCE CHARACTERISTICS:
50 (A) LENGTH: 8083 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: double
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- 55 (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
- 60 (vi) ORIGINAL SOURCE:
(A) ORGANISM: Bovine viral diarrhea virus
(B) STRAIN: 2724
(C) INDIVIDUAL ISOLATE: pBEVtkex-3::p53
- 65 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

	GCGCCCAATA CGCAAACCGC CTCTCCCCGC GCGTTGGCCG ATTCATTAAT GCAGCTGGCA	60
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5	CACTCATTAG GCACCCCAGG CTTTACACTT TATGCTTCCG GCTCGTATGT TGTGTGGAAT	180
	TGTGAGCGGA TAACAATTTC ACACAGGAAA CAGCTATGAC CATGATTACG CCAAGCTAGC	240
	TTGCATGCCT GCAGGTCGAC TTCCGCGCCC GCGGCGTCTG CCTTCGCCAG CAGGTTGTCC	300
10	GCGGCCGCTG CCGGCCTGGT TCCGCGCCCG CCGCCTCGCG GCCAGCTCCC GCGCGGGCGC	360
	GTCCGCGTCC CCAACTCCGC GCGAAGACGG GCTCGTCCCA GAAGCGCAGC GGAAAGGCCG	420
15	GCGTATAAAA TTTCGCTCGT CCGGTACAAA GACGCGGTCC GCGACTGCGT GGATGTCCAC	480
	GCCCAGGCAA GCAAACCTA AACGCCCAG CCGCATGGCC CCGATGCCGC CACAAAGAGC	540
	GCCGAAATTT CGCCCAGGCA CGCCGCGCCG CCCGACGCGT CTTTAGCGCA CCCGCCGGCG	600
20	CTGTTGCCCG CGTGCTGCT GGCCGCCCAC CGGCGGCCG TGTCCCCGGC CTCAGCAGGG	660
	CCGGGGTCTGC CGGCGGGCGG CCGCGGGGTG CGGCCACAGC CGCCCTTTTG CCCGTAGCCA	720
25	GGGGAAGCGG CTGCCCCCTT TGCCGCGCGG GCCGCGGTTG CTCGGCTTTG CGTTTGCCCC	780
	GCGGCGATCG CCCCCTCGC CGCGAACGCG CGCGCGCGAA TGGGGCGTAC TCGGCGAGCC	840
	CGGCTATTAT AGCCTCAAGG CGCGCCGCGT TGCTAGCGAT CGTCTGGGCC GGCAGGCGCG	900
30	TCACTCTGAG CACGCGCATG CCCCCTGGG AGACGAACAC CTGCACCGGC GCTAGGACCA	960
	CCGGGTCTGG GCCCGGGGGG GCGAGATCGC GCACAAGCCG GGCCGAGTCG CGCAGCTGCC	1020
35	GCAGCCCCC GAGGCGCTGG TCCATCTTGC TGGGCGTGTT CATGTTCTGT GAAAAACGGC	1080
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40	CCACGACCTG CTTAAAAACC TCCGGGGCGC CAACTTGCC CAAAAGCTGG GCGAGGCGCG	1260
	GGCGCAGCTT CTGCGCGCCA ACCGCCGCGC GTGCGTCGCA AGCCAGCGCC TCGTAAAAGC	1320
45	GGCTGTGGCA CCGGATCCCG GCGCGCAGGC GCGCACGTCG GTCGCGGTCG CGCGCCATGG	1380
	CCGAGCCCGC GCGCGCTCTC CGCGTCGTGC GTATCTACCT GGACGGCGCG CACGGGCAGG	1440
	GAAAGACAAC AACGGGCCGC GCGCTCGCGG CCGCTTCCAC CGCTGGGGAG GGCCTGCTCT	1500
50	TTTTCCCGGA GCCGATGGCG TACTGGCGCA CGATGTTTGG TACGGACGCC TTAAGTGGGA	1560
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55	CCGGCGCACC GCGCAGACGC GGACGCGCG GGCCTGGTTG CGTACTACCA GGCCAGGTTT	1680
	GCGGCCCCGT ACTTAATTTT GCACGCGCGT GTCCGCGCTG CTGCGCCGCC TGGGCCGGCG	1740
	CCGGGCGGCG AGCTGGTGGG CCCTCGTGTT CGACCGCCAC CCCGTGGCGC GCGTGCCTCT	1800
60	GCTACCCCTT CGCCCGCTAC TGCCCTCCCG AGATCAACGC GGAAGATCCG AATTCCTCGA	1860
	CCTGCAGTGA ATAATAAAAT GTGTGTTTGT CCGAAATACG CGTTTGAGAT TTCTGTCCCG	1920
65	ACTAAATTCG TGTGCGCGA TAGTGGTGTT TATCGCCGAT AGAGATGGCG ATATTGAAA	1980
	AATCGATATT TGAAATATG GCATATTGAA AATGTCGCCG ATGTGAGTTT CTGTGTAAC	2040

	GATATCGCCA	TTTTTCCAAA	AGTTGATTTT	TGGGCATACG	CGATATCTGG	CGATACGCTT	2100
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5	CAAATATCGC	AGTTTCGATA	TAGGTGACAG	ACGATATGAG	GCTATATCGC	CGATAGAGGC	2220
	GACATCAAGC	TGGCAGATGG	CCAATGCATA	TCGATCTATA	CATTGAATCA	ATATTGGCCA	2280
10	TTAGCCATAT	TATTCATTGG	TTATATAGCA	TAAATCAATA	TTGGCTATTG	GCCATTGCAT	2340
	ACGTTGTATC	CATATCATAA	TATGTACATT	TATATTGGCT	CATGTCCAAC	ATTACCGCCA	2400
	TGTTGACATT	GATTATTGAC	TAGTTATTAA	TAGTAATCAA	TTACGGGGTC	ATTAGTTCAT	2460
15	AGCCCATATA	TGGAGTTCCG	CGTTACATAA	CTTACGGTAA	ATGGCCCCGC	TGGCTGACCG	2520
	CCCAACGACC	CCCGCCCAT	GACGTCAATA	ATGACGTATG	TTCCCATAGT	AACGCCAATA	2580
20	GGGACTTTCC	ATTGACGTCA	ATGGGTGGAG	TATTTACGGT	AAACTGCCCC	CTTGCCAGTA	2640
	CATCAAGTGT	ATCATATGCC	AAGTACGCCC	CCTATTGACG	TCAATGACGG	TAAATGGCCC	2700
	GCCTGGCATT	ATGCCCAGTA	CATGACCTTA	TGGGACTTTC	CTACTTGGCA	GTACATCTAC	2760
25	GTATTAGTCA	TCGCTATTAC	CATGGTGATG	CGGTTTTGGC	AGTACATCAA	TGGGCGTGGA	2820
	TAGCGGTTTG	ACTCACGGGG	ATTTCCAAGT	CTCCACCCCA	TTGACGTCAA	TGGGAGTTTG	2880
30	TTTTGGCACC	AAAATCAACG	GGACTTTCCA	AAATGTCGTA	ACAACTCCGC	CCCATTGACG	2940
	CAAATGGGCG	GTAGGCGTGT	ACGGTGGGAG	GTCTATATAA	GCAGAGCTCG	TTTAGTGAAC	3000
	CGTCAGATCG	CCTGGAGACG	CCATCCACGC	TGTTTTGACC	TCCATAGAAG	ACACCGGGAC	3060
35	CGATCCAGCC	TCCGCGGCAA	GCTGATCCGT	CAGGGGCCAG	ATGGTACAGG	GCATCCTATG	3120
	GCTACTACTG	ATAACAGGGG	TACAAGGGGA	CATTGACTGC	AAACCTGAAC	ACTCATACGC	3180
40	CATAGCCAGG	AATGATAGAA	TTGGCCCATT	AGGAGCTGAA	GGCCTCACCA	CTGTTTGGAA	3240
	GGATTACTCA	CATGAAATGA	AGCTGGAAGA	CACAATGGTC	ATAGCTTGGT	GCAAAGACGG	3300
	TAAGTTTACA	TACCTCTCAA	GGTGCACAAG	AGAACTAGA	TATCTTGCAA	TTCTGCATTC	3360
45	AAGAGCCTTG	CAGACCAGTG	TGGTATTCAA	AAAACTTTTT	GAGGGGCAA	GGCAAGGGGA	3420
	AACATTTGAA	ATGGCTGACG	ACTTTGAATT	TGGACTCTGC	CCATGCGATG	CCAATCCCGT	3480
50	AGTAAGAGGG	AAGTTCAATA	CAACACTGCT	AAACGGACCG	GCCTTCCAGA	TGGTATGCCC	3540
	TATAGGATGG	ACAGGAACTG	TGAGCTGTAT	GTTAGCTAAT	AGGGACACCC	TAGACACAGC	3600
	AGTAGTGCCT	GTGTATAAGA	GGTCCAAACC	ATTCCCTTAT	AGACAAGGTT	GTATCACCCA	3660
55	AAGAACTCTG	GGGGAGGATC	TCTATAACTG	TGATCTTGGA	GGGAATTGGA	CTGTGTGTGAC	3720
	TGGGGACCAG	CTACAATACA	CAGGAGGCCC	TGTCGAATCT	TGCAAGTGGT	GTGGTTATAA	3780
60	ATTCCAAAAA	AGTGAGGGGT	TGCCACACTA	CCCCATCGGC	AAGTGAGGT	TGAAGAATGA	3840
	GACTGGCTAC	AGATTGTAG	ACGGCACCAC	TTGCAACAGA	GAGGGTGTAG	CCATAGTACC	3900
	ACAAGGATTG	GTAAAGTGTA	AGATAGGAGA	CACAATCGTA	CAGGTCATAG	CTCTTGACAC	3960
65	CAAACTTGGG	CCTATGCCTT	GCAAGCCATA	TGAGATCATA	CCAAGTGAGG	GGCCTGTAGA	4020
	AAAGACGGCA	TGCACCTTCA	ACTACACGAG	GACATTAAAA	AATAAATATT	TTGAGCCCAG	4080

	AGACAGTTAC TTCCAGCAAT ACATGCTAAA AGGAGATTAT CAATACTGGT TCGACCTGGA	4140
	GGTCACTGAC CATCATCGGG ATTACTTCGC CGAGTCCATA TTGGTGGTGG TGGTAGCTTT	4200
5	ACTGGGTGGA AGATACGTGC TCTGGTTACT GGTAACATAC ATGGTCCTAT CAGAACAAAA	4260
	GGCCTTGGGG ACCCAATATG GGGCAGGGGA AGTGGTGATG ATGGGTAAC TGTAAACACA	4320
	TGACAGTATT GAAGTGGTGA CATATTTCTT GTTGTTATAC CTAAGTCTAA GAGAGGAGGC	4380
10	TGTAAAGAAG TGGGTCTTAC TCTTATACCA CCTTGATTGA TTGAGGATCA GCTTATCCAG	4440
	GGTCGACCTC AGGCATGCAA GCTCAGATCC GCTGTGCCTT CTAGTTGCCA GCCATCTGTT	4500
15	GTTTGCCCCCT CCCCCGTGCC TTCCTTGACC CTGGAAGGTG CCACTCCCAC TGTCTTTTCC	4560
	TAATAAAATG AGGAAATTGC ATCGCATTGT CTGAGTAGGT GTCATTCTAT TCTGGGGGGT	4620
	GGGGTGGGGC AGGACAGCAA GGGGGAGGAT TGGGAAGACA ATAGCAGGCA TGCTGGGGAT	4680
20	GCGGTGGGCT CTATGGGTAC CCAGGTGCTG AAGAATTGAC CCGGTTCCCTC CTGGGCCAGA	4740
	AAGAAGCAGG CACATCCCCT TCTCTGTGAC ACACCCTGTC CACGCCCCTG GTTCTTAGTT	4800
25	CCAGCCCCAC TCATAGGACA CTCATAGCTC AGGAGGGCTC CGCTTCAATC CCACCCGCTA	4860
	AAGTACTTGG AGCGGTCTCT CCCTCCCTCA TCAGCCCACC AAACCAAACC TAGCCTCCAA	4920
	GAGTGGGAAG AAATTAAAGC AAGATAGGCT ATTAAGTGCA GAGGGAGAGA AAATGCCTCC	4980
30	AACATGTGAG GAAGTAATGA TAGAAATCAT AGAATTGAGA TCTCGAGGTG TTCGTGCTGG	5040
	ACGTGTCCGC GCGCCAGAC GCGTGCGCGG CCGCCGTACT GGACATGCGG CCCGCCATGC	5100
35	AGGCCGCTTG CCGCGACGGG GCGGCGGGCG CGACGCTGGC GACCCTGGCG CGTCAGTTCC	5160
	CGCTAGAGAT GCGGGGGGAG GCCACGGCGG GCCCTAGGGG ACTATAAAGC TGCCCCCTGCG	5220
	CTCGCTCGCT CGCTGCATTT GCGCCCCGAT CGCCTTACGG GGAATCGGCG CTCGGCGGAT	5280
40	CCCCCCCCG CCGCGCCGCG AAGCAGGCCG CCAGACAAAA AAATGCGGCG CCCGCTCTGC	5340
	GCGGCGCTAT TGGCAGCGGC TGTCTCGCG CTCGCGCGCG GCGCCCCCGC CGCCGCCCCG	5400
45	GGCGGGGGCG CCGAAGCCAG GGCAGCACAG AGACGCCCCG TACGAAATCG AAGAGTGGGA	5460
	AATGGTGGTC GGAGCCGGGC CGGCCGTGCA CACGTTTACC ATCCGCTGCC TCGGGCCGCG	5520
	GGGCATTGAG CGCGTGGCCC ACATTGCAAA CCTCAGCCGG CTGCTGGACG GGTACATAGC	5580
50	GGTCCACGTT GACGTTGCGC GCACCTCTGG CCTGCGGGAC GCCATGTTTT TCCTGCCGCG	5640
	CGCGGCCGTC GACTCTAGAG GATCCCCGGG TACCGAGCTC GAATTCAGTG GCCGTCGTTT	5700
55	TACAACGTCG TGAAGTGGAA AACCTGGCG TTACCCAACT TAATCGCCTT GCAGCACATC	5760
	CCCCCTTCGC CAGCTGGCGT AATAGCGAAG AGGCCCGCAC CGATCGCCCT TCCCAACAGT	5820
	TGCGCAGCCT GAATGGCGAA TGGCGCCTGA TGCGGTATTT TCTCCTTACG CATCTGTGCG	5880
60	GTATTTTACA CCGCATATGG TGCACTCTCA GTACAATCTG CTCTGATGCC GCATAGTTAA	5940
	GCCAGCCCCG ACACCCGCCA ACACCCGCTG ACGCGCCCTG ACGGGCTTGT CTGCTCCCCG	6000
65	CATCCGCTTA CAGACAAGCT GTGACCGTCT CCGGGAGCTG CATGTGTCAG AGGTTTTTAC	6060
	CGTCATCACC GAAACGCGCG AGACGAAAGG GCCTCGTGAT ACGCCTATTT TTATAGGTTA	6120

	ATGTCATGAT AATAATGGTT TCTTAGACGT CAGGTGGCAC TTTTCGGGGA AATGTGCGCG	6180
	GAACCCCTAT TTGTTTATTT TTCTAAATAC ATTCAAATAT GTATCCGCTC ATGAGACAAT	6240
5	AACCCTGATA AATGCTTCAA TAATATTGAA AAAGGAAGAG TATGAGTATT CAACATTTCC	6300
	GTGTCGCCCT TATTCCTTT TTTGCGGCAT TTTGCCTTCC TGTTTTTGCT CACCCAGAAA	6360
	CGCTGGTGAA AGTAAAAGAT GCTGAAGATC AGTTGGGTGC ACGAGTGGGT TACATCGAAC	6420
10	TGGATCTCAA CAGCGGTAAG ATCCTTGAGA GTTTTCGCCC CGAAGAACGT TTTCCAATGA	6480
	TGAGCACTTT TAAAGTTCTG CTATGTGGCG CGGTATTATC CCGTATTGAC GCCGGGCAAG	6540
15	AGCAACTCGG TCGCCGCATA CACTATTCTC AGAATGACTT GGTGAGTAC TCACCAGTCA	6600
	CAGAAAAGCA TCTTACGGAT GGCATGACAG TAAGAGAATT ATGCAGTGCT GCCATAACCA	6660
	TGAGTGATAA CACTGCGGCC AACTTACTTC TGACAACGAT CGGAGGACCG AAGGAGCTAA	6720
20	CCGCTTTTTT GCACAACATG GGGGATCATG TAACTCGCCT TGATCGTTGG GAACCGGAGC	6780
	TGAATGAAGC CATACCAAAC GACGAGCGTG ACACCACGAT GCCTGTAGCA ATGGCAACAA	6840
25	CGTTGCGCAA ACTATTAAC T GCGGAAC T TACTCTAGC TTCCCGGCAA CAATTAATAG	6900
	ACTGGATGGA GCGGATAAA GTTGACAGG CACTTCTGCG CTCGGCCCTT CCGGCTGGCT	6960
	GGTTTATTGC TGATAAATCT GGAGCCGGTG AGCGTGGGTC TCGCGGTATC ATTGCAGCAC	7020
30	TGGGGCCAGA TGGTAAGCCC TCCCGTATCG TAGTTATCTA CACGACGGGG AGTCAGGCAA	7080
	CTATGGATGA ACGAAATAGA CAGATCGCTG AGATAGGTGC CTCACTGATT AAGCATTGGT	7140
35	AACTGTCAGA CCAAGTTTAC TCATATATAC TTTAGATTGA TTTAAAACTT CATTTTTAAT	7200
	TTAAAAGGAT CTAGGTGAAG ATCCTTTTTG ATAATCTCAT GACCAAAATC CCTTAACGTG	7260
	AGTTTTTCGT CCACTGAGCG TCAGACCCCG TAGAAAAGAT CAAAGGATCT TCTTGAGATC	7320
40	CTTTTTTCT GCGCGTAATC TGCTGCTTGC AAACAAAAA ACCACCGCTA CCAGCGGTGG	7380
	TTTGTTCGCC GGATCAAGAG CTACCAACTC TTTTCCGAA GGTAAC TGC TTCAGCAGAG	7440
45	CGCAGATACC AAATACTGTC CTTCTAGTGT AGCCGTAGTT AGGCCACCAC TTCAAGAACT	7500
	CTGTAGCACC GCCTACATAC CTCGCTCTGC TAATCCTGTT ACCAGTGGCT GCTGCCAGTG	7560
	GCGATAAGTC GTGTCTTACC GGGTTGGACT CAAGACGATA GTTACCGGAT AAGGCGCAGC	7620
50	GGTCGGGCTG AACGGGGGGT TCGTGACAC AGCCAGCTT GGAGCGAACG ACCTACACCG	7680
	AACTGAGATA CCTACAGCGT GAGCTATGAG AAAGCGCCAC GCTTCCCGAA GGGAGAAAGG	7740
55	CGGACAGGTA TCCGTAAGC GGCAGGGTCG GAACAGGAGA GCGCACGAGG GAGCTTCCAG	7800
	GGGGAAACGC CTGGTATCTT TATAGTCCTG TCGGGTTTCG CCACCTCTGA CTGAGCGTC	7860
	GATTTTTGTG ATGCTCGTCA GGGGGGCGGA GCCTATGGAA AAACGCCAGC AACGCGGCCT	7920
60	TTTACGGTT CCTGGCCTTT TGCTGGCCTT TTGCTCACAT GTTCTTTCCT GCGTTATCCC	7980
	CTGATTCTGT GGATAACCGT ATTACCGCCT TTGAGTGAGC TGATAACGCT CGCCGAGCC	8040
65	GAACGACCGA GCGCAGCGAG TCAGTGAGCG AGGAAGCGGA AGA	8083

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 8149 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
- (v) ORIGINAL SOURCE:
 (A) ORGANISM: Bovine viral diarrhea virus
 (B) STRAIN: 2724
 (C) INDIVIDUAL ISOLATE: pBHVtkex-1::gBGH/p53

20

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

	GCGCCCAATA CGCAAACCGC CTCTCCCCGC GCGTTGGCCG ATTCATTAAT GCAGCTGGCA	60
25	CGACAGGTTT CCCGACTGGA AAGCGGGCAG TGAGCGCAAC GCAATTAATG TGAGTTAGCT	120
	CACTCATTAG GCACCCCAGG CTTTACACTT TATGCTTCCG GCTCGTATGT TGTGTGGAAT	180
	TGTGAGCGGA TAACAATTTC ACACAGGAAA CAGCTATGAC CATGATTACG CCAAGCTAGC	240
30	TTGCATGCCT GCAGGTCGAC TTCCGCGCCC GCGGCGTCTG CCTTCGCCAG CAGGTTGTCC	300
	GCGGCCGCTG CCGGCCTGGT TCCGCGCCCG CCGCCTCGCG GCCAGCTCCC GCGCGGGCGC	360
35	GTCCGCGTCC CCAACTCCGC GCGAAGACGG GCTCGTCCCA GAAGCGCAGC GGAAAGGCCG	420
	GCGTATAAAA TTTCGCTCGT CCGGTACAAA GACGCGGTCC GCGACTGCGT GGATGTCCAC	480
	GCCCAGGCAA GCAAACCTTA AACGCCCAG CCGCATGGCC CCGATGCCGC CACAAAGAGC	540
40	GCCGAAATTT CGCCCAGGCA CGCCGCGCCG CCCGACGCGT CTTTAGCGCA CCCGCCGGCG	600
	CTGTTGCCCG CGTGCCCTGCT GGCCGCCCAC CGGCGGGCCG TGTCCCCGGC CTCAGCAGGG	660
45	CCGGGGTCGC CGGCGGGCGG CCGCGGGGTG CGGCCACAGC CGCCCTTTTG CCCGTAGCCA	720
	GGGGAAGCGG CTGCCCCCTC TGCCGCCGCG GCCGCGGTTG CTCGGCTTTG CGTTTGCCCC	780
	GCGGCGATCG CCCCGCTCGC CGCGAACGCG CGCGCGCGAA TGGGGCGTAC TCGGCGAGCC	840
50	CGGCTATTAT AGCCTCAAGG CGCGCCGCGT TGCTAGCGAT CGTCTGGGCC GGCAGGCGCG	900
	TCACTCTGAG CACGCGCATG CCCCGCTGGG AGACGAACAC CTGCACCGGC GCTAGGACCA	960
55	CCGGGTCTGG GCCCGGGGGG GCGAGATCGC GCACAAGCCG GGCCGAGTCG CGCAGCTGCC	1020
	GCAGCCCCCC GAGGCGCTGG TCCATCTTGC TGGGCGTGTT CATGTTCTGT GAAAAACGGC	1080
	ACGTCTTCAG CTCCACGATA AGACAGACGG CCCGGGCGTG CCCTGCCTCC GCGACCCGGA	1140
60	GTAGGCACAC GCAATCGGGC CGCCGGCTTT GCAGGTTTAC CTCAAAGCTC AGAGACACGC	1200
	CCACGACCTG CTTAAAAACC TCCGGGGCGC CAAACTTGCC CAAAAGCTGG GCGAGGCGCG	1260
65	GGCGCAGCTT TTGCGCGCCA ACCGCGCGC GTGCGTCGCA AGCCAGCGCC TCGTAAAAGC	1320
	GGCTGTGGCA CCGGATCCCG GCGCGCAGGC GCGCACGTCG GTCGCGGTCTG CGCGCCATGG	1380

	CCGAGCCCGC	GCGCGCTCTC	CGCGTCGTGC	GTATCTACCT	GGACGGCGCG	CACGGGCAGG	1440
	GAAAGACAAC	AACGGGCCCGC	GCGCTCGCGG	CCGCTTCCAC	CGCTGGGGAG	GGCGTGCTCT	1500
5	TTTTCCCGGA	GCCGATGGCG	TACTGGCGCA	CGATGTTTGG	TACGGACGCC	TTAAGTGGGA	1560
	TCCTCGCGGC	GTCTGCGCGA	TGCGCCGCAG	CCTCGCACGG	GAGCGCACGC	GCGCGGCGGG	1620
	CCGGCGCACC	GCGCAGACGC	GGACGCGGCG	GGCCTGGTTG	CGTACTACCA	GGCCAGGTTT	1680
10	GCGGCCCCGT	ACTTAATTTT	GCACGCGCGT	GTCCGCGCTG	CTGCGCCGCC	TGGGCCGGCG	1740
	CCGGGCGGCG	AGCTGGTGGA	CCCTCGTGTT	CGACCGCCAC	CCCGTGGCGC	GCGTGCTCT	1800
15	GCTACCCCTT	CGCCCGCTAC	TGCCTCCGCG	AGATCAACGC	GGAAGATCTC	AATTCTATGA	1860
	TTTCTATCAT	TACTTCCTCA	CATGTTGGAG	GCATTTTCTC	TCCCTCTGCA	CTTAATAGCC	1920
	TATCTTGCTT	TAATTTCTTC	CCACTCTTGG	AGGCTAGGTT	TGGTTTGGTG	GGCTGATGAG	1980
20	GGAGGGAGAG	ACCGCTCCAA	GTACTTTAGC	GGGTGGGATT	GAAGCGGAGC	CCTCCTGAGC	2040
	TATGAGTGTC	CTATGAGTGG	GGCTGGAAct	AAGAACCAGG	GGCGTGGACA	GGGTGTGTCA	2100
25	CAGAGAAGGG	GATGTGCCTG	CTTCTTTCTG	GCCCAGGAGG	AACCGGGTCA	ATTCTTCAGC	2160
	ACCTGGGTAC	CCATAGAGCC	CACCGCATCC	CCAGCATGCC	TGCTATTGTC	TTCCCAATCC	2220
	TCCCCCTTGC	TGCTCTGCCC	CACCCACCCC	CCCAGAATAG	AATGACACCT	ACTCAGACAA	2280
30	TGCGATGCAA	TTTCCTCATT	TTATTAGGAA	AGGACAGTGG	GAGTGGCACC	TTCCAGGGTC	2340
	AAGGAAGGCA	CGGGGGAGGG	GCAAACAACA	GATGGCTGGC	AACTAGAAGG	CACAGCGGAT	2400
35	CTGAGCTTGC	ATGCCTGAGG	TCGACCCTGG	ATAAGCTGAT	CCTCAATCAA	TCAAGGTGGT	2460
	ATAAGAGTAA	GACCCACTTC	TTACAGCCT	CCTCTCTTAG	CAGTAGGTAT	AACAACAAGA	2520
	AATATGTCAC	CACTTCAATA	CTGTCAATG	TTAGCAAGTT	ACCCATCATC	ACCACTTCCC	2580
40	CTGCCCCATA	TTGGGTCCCC	AAGGCCTTTT	GTTCTGATAG	GACCATGTAT	GTTACCAGTA	2640
	ACCAGAGCAC	GTATCTTCCA	CCCAGTAAAG	CTACCACCAC	CACCAATATG	GACTCGGCGA	2700
45	AGTAATCCCG	ATGATGGTCA	GTGACCTCCA	GGTCGAACCA	GTATTGATAA	TCTCCTTTTA	2760
	GCATGTATTG	CTGGAAGTAA	CTGTCTCTGG	GCTCAAAATA	TTTATTTTTT	AATGTCTCTG	2820
	TGTAGTTGAA	GGTGCATGCC	GTCTTTTCTA	CAGGCCCTCT	ACTTGGTATG	ATCTCATATG	2880
50	GCTTGCAAGG	CATAGGCCCA	AGTTTGGTGT	CAAGAGCTAT	GACCTGTACG	ATTGTGTCTC	2940
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55	TGCCGTCTAC	AAATCTGTAG	CCAGTCTCAT	TCTTCAACCT	ACACTTGCCG	ATGGGGTAGT	3060
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	CTCCTGTGTA	TTGTAGCTGG	TCCCCAGTCA	CACAAGTCCA	ATTCCCTCCA	AGATCACAGT	3180
60	TATAGAGATC	CTCCCCCAGA	GTTCTTTGGG	TGATACAACC	TTGTCTATAA	GGGAATGGTT	3240
	TGGACCTCTT	ATACACACGC	ACTACTGCTG	TGTCTAGGGT	GTCCCTATTA	GCTAACATAC	3300
65	AGCTCACAGT	TCCTGTCCAT	CCTATAGGGC	ATACCATCTG	GAAGGCCGGT	CCGTTTAGCA	3360
	GTGTTGTATT	GAACCTCCCT	CTTACTACGG	GATTGGCATC	GCATGGGCAG	AGTCCAAATT	3420

	CAAAGTCGTC AGCCATTTCA AATGTTTCCC CTTGCCTTTG CCCCTCGAAA AGTTTTTTTGA	3480
	ATACCACACT GGTCTGCAAG GCTCTTGAAT GCAGAATTGC AAGATATCTA GTTCTCTTTG	3540
5	TGCACCTTGA GAGGTATGTA AACTTACCGT CTTTGACCA AGCTATGACC ATTGTGTCTT	3600
	CCAGCTTCAT TTCATGTGAG TAATCCTTCC AAACAGTGGT GAGGCCCTCA GCTCCTAATG	3660
10	GGCCAATTCT ATCATTCCCTG GCTATGGCGT ATGAGTGTTT AGGTTTGCAG TCAATGTCCC	3720
	CTTGTACCCC TGTATCAGT AGTAGCCATA GGATCCCTGG GAAGGCGCCC ACCACCTGAG	3780
	TCCAGGGCAG GCAGAGCAGG GCGAAAGCCA GGAGCAGGGA GGTCCGGGGG CCTGCAGCCA	3840
15	TCATGTCGAA GCTTGCCGCG GAGGCTGGAT CGGTCCCGGT GTCTTCTATG GAGGTCAAAA	3900
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	ACCTCCCACC GTACACGCCT ACCGCCCAT TGCCTCAATG GGGCGGAGTT GTTACGACAT	4020
20	TTTGGAAAGT CCCGTTGATT TTGGTGCCAA AACAACTCC CATTGACGTC AATGGGGTGG	4080
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25	CGCATACCA TGGTAATAGC GATGACTAAT ACGTAGATGT ACTGCCAAGT AGGAAAGTCC	4200
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30	TACTCCACCC ATTGACGTCA ATGGAAAGTC CCTATTGGCG TTACTATGGG AACATACGTC	4380
	ATTATTGACG TCAATGGGCG GGGGTCGTTG GCGGGTCAGC CAGGCGGGCC ATTTACCGTA	4440
35	AGTTATGTAA CGCGGAATC CATATATGGG CTATGAACTA ATGACCCCGT AATTGATTAC	4500
	TATTAATAAC TAGTCAATAA TCAATGTCAA CATGGCGGTA ATGTTGGACA TGAGCCAATA	4560
	TAAATGTACA TATTATGATA TGGATACAAC GTATGCAATG GCCAATAGCC AATATTGATT	4620
40	TATGCTATAT AACCAATGAA TAATATGGCT AATGGCCAAT ATTGATTCAA TGTATAGATC	4680
	GATATGCATT GGCCATGTGC CAGCTTGATG TCGCCTCTAT CGGCATATA GCCTCATATC	4740
45	GTCTGTCACC TATATCGAAA CTGCGATATT TCGGACACAC AGAATCGCCC AAGTCACCAA	4800
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	CAAAAATCAA CTTTTGAAA AATGGCGATA TCAGTTACAC AGAACTCAC ATCGGCGACA	4920
50	TTTTCAATAT GCCATATTTT CAAATATCGA TTTTCCAAT ATCGCCATCT CTATCGGCGA	4980
	TAAACACCAC TATCGCGCGA CATGAATTTA GTCGGGACAG AAATCTCAA CGCGTATTTT	5040
55	GGACAAACAC ACATTTTATT ATTCACTGCA GGTCGAGGAA TTCGGATCTC GAGGTGTTTCG	5100
	TGCTGGACGT GTCCGCGGCG CCAGACGCGT GCGCGGCCGC CGTACTGGAC ATGCGGCCCG	5160
	CCATGCAGGC CGCTTGCGCG GACGGGGCGG CGGGCGCGAC GCTGGCGACC CTGGCGCGTC	5220
60	AGTTCGCGCT AGAGATGGCG GGGGAGGCCA CGGCGGGCCC TAGGGGACTA TAAAGCTGCC	5280
	CCTGCGCTCG CTCGCTCGCT GCATTTGCGC CCCGATCGCC TTACGGGGAC TCGGCGCTCG	5340
65	GCGGATCCCC TCCCGGCCCG GCCGCGAAGC AGGCCGCCAG ACAAAAAAAT GCGGCGCCCG	5400
	CTCTGCGCGG CGCTATTGGC AGCGGCTGTC CTCGCGCTCG CCGCGGGCGC CCGCGCGGCC	5460

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5	GCCGCGGGGC	ATTGAGCGCG	TGGCCCACAT	TGCAAACCTC	AGCCGGCTGC	TGGACGGGTA	5640
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	TCGTTTTACA	ACGTCGTGAC	TGGGAAAACC	CTGGCGTTAC	CCAACTTAAT	CGCCTTGCAG	5820
	CACATCCCCC	TTTCGCCAGC	TGGCGTAATA	GCGAAGAGGC	CCGCACCGAT	CGCCCTTCCC	5880
15	AACAGTTGCG	CAGCCTGAAT	GGCGAATGGC	GCCTGATGCG	GTATTTTCTC	CTTACGCATC	5940
	TGTGCGGTAT	TTACACCCGC	ATATGGTGCA	CTCTCAGTAC	AATCTGCTCT	GATGCCGCAT	6000
20	AGTTAAGCCA	GCCCCGACAC	CCGCCAACAC	CCGCTGACGC	GCCCTGACGG	GCTTGTCTGC	6060
	TCCCGGCATC	CGCTTACAGA	CAAGCTGTGA	CCGTCTCCGG	GAGCTGCATG	TGTCAGAGGT	6120
	TTTCACCGTC	ATCACCGAAA	CGCGCGAGAC	GAAAGGGCCT	CGTGATACGC	CTATTTTTAT	6180
25	AGGTAAATGT	CATGATAATA	ATGGTTTCTT	AGACGTCAGG	TGGCACTTTT	CGGGGAAATG	6240
	TGCGCGGAAC	CCCTATTTGT	TTATTTTTCT	AAATACATTC	AAATATGTAT	CCGCTCATGA	6300
30	GACAATAACC	CTGATAAATG	CTTCAATAAT	ATTGAAAAAG	GAAGAGTATG	AGTATTCAAC	6360
	ATTTCCGTGT	CGCCCTTATT	CCCTTTTTTG	CGGCATTTTG	CCTTCCTGTT	TTTGCTCACC	6420
	CAGAAACGCT	GGTGAAAAGTA	AAAGATGCTG	AAGATCAGTT	GGGTGCACGA	GTGGGTTACA	6480
35	TCGAACTGGA	TCTCAACAGC	GGTAAGATCC	TTGAGAGTTT	TCGCCCCGAA	GAACGTTTTT	6540
	CAATGATGAG	CACTTTTAAA	GTTCTGCTAT	GTGGCGCGGT	ATTATCCCGT	ATTGACGCCG	6600
40	GGCAAGAGCA	ACTCGGTCGC	CGCATACACT	ATTCTCAGAA	TGACTTGGTT	GAGTACTCAC	6660
	CAGTCACAGA	AAAGCATCTT	ACGGATGGCA	TGACAGTAAG	AGAATTATGC	AGTGCTGCCA	6720
	TAACCATGAG	TGATAACACT	GCGGCCAACT	TACTTCTGAC	AACGATCGGA	GGACCGAAGG	6780
45	AGCTAACCGC	TTTTTTGCAC	AACATGGGGG	ATCATGTAAC	TCGCCTTGAT	CGTTGGGAAC	6840
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50	CAACAACGTT	GCGCAAACCT	TTAACTGGCG	AACTACTTAC	TCTAGCTTCC	CGGCAACAAT	6960
	TAATAGACTG	GATGGAGGCG	GATAAAGTTG	CAGGACCACT	TCTGCGCTCG	GCCCTTCCGG	7020
	CTGGCTGGTT	TATTGCTGAT	AAATCTGGAG	CCGGTGAGCG	TGGGTCTCGC	GGTATCATTG	7080
55	CAGCACTGGG	GCCAGATGGT	AAGCCCTCCC	GTATCGTAGT	TATCTACACG	ACGGGGAGTC	7140
	AGGCAACTAT	GGATGAACGA	AATAGACAGA	TCGCTGAGAT	AGGTGCCTCA	CTGATTAAGC	7200
60	ATTGGTAACT	GTCAGACCAA	GTTTACTCAT	ATATACTTTA	GATTGATTTA	AAACTTCATT	7260
	TTTAATTTAA	AAGGATCTAG	GTGAAGATCC	TTTTTGATAA	TCTCATGACC	AAAATCCCTT	7320
	AACGTGAGTT	TTCGTTCCAC	TGAGCGTCAG	ACCCCGTAGA	AAAGATCAAA	GGATCTTCTT	7380
65	GAGATCCTTT	TTTTCTGCGC	GTAATCTGCT	GCTTGCAAAC	AAAAAAACCA	CCGCTACCAG	7440
	CGGTGGTTTG	TTTGCCGGAT	CAAGAGCTAC	CAACTCTTTT	TCCGAAGGTA	ACTGGCTTCA	7500

GCAGAGCGCA GATACCAAAT ACTGTCCTTC TAGTGTAGCC GTAGTTAGGC CACCACTTCA 7560
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 5 CCAGTGGCGA TAAGTCGTGT CTTACCGGGT TGGACTCAAG ACGATAGTTA CCGGATAAGG 7680
 CGCAGCGGTC GGGCTGAACG GGGGGTTCGT GCACACAGCC CAGCTTGGAG CGAACGACCT 7740
 ACACCGAACT GAGATACCTA CAGCGTGAGC TATGAGAAAG CGCCACGCTT CCCGAAGGGA 7800
 10 GAAAGGCGGA CAGGTATCCG GTAAGCGGCA GGGTCGGAAC AGGAGAGCGC ACGAGGGAGC 7860
 TTCCAGGGGG AAACGCCTGG TATCTTTATA GTCCTGTCGG GTTTCGCCAC CTCTGACTTG 7920
 15 AGCGTCGATT TTTGTGATGC TCGTCAGGGG GCGGAGCCT ATGGA AAAAC GCCAGCAACG 7980
 CGGCCTTTTT ACGGTTCCTG GCCTTTTGCT GGCCTTTTGC TCACATGTTT TTTCTGCGT 8040
 TATCCCCTGA TTCTGTGGAT AACCGTATTA CCGCCTTTGA GTGAGCTGAT ACCGCTCGCC 8100
 20 GCAGCCGAAC GACCGAGCGC AGCGAGTCAG TGAGCGAGGA AGCGGAAGA 8149

(2) INFORMATION FOR SEQ ID NO:3:

- 25 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 8135 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear
- 30 (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- 35 (iv) ANTI-SENSE: NO
- (vi) ORIGINAL SOURCE:
 (A) ORGANISM: Bovine viral diarrhea virus
 (B) STRAIN: 2724
 40 (C) INDIVIDUAL ISOLATE: pBHvtkex-1::gIII/p53

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

45 GCGCCCAATA CGCAAACCGC CTCTCCCCGC GCGTTGGCCG ATTCATTAAT GCAGCTGGCA 60
 CGACAGGTTT CCCGACTGGA AAGCGGGCAG TGAGCGCAAC GCAATTAATG TGAGTTAGCT 120
 CACTCATTAG GCACCCAGG CTTTACACTT TATGCTTCCG GCTCGTATGT TGTGTGGAAT 180
 50 TGTGAGCGGA TAACAATTTC ACACAGGAAA CAGCTATGAC CATGATTACG CCAAGCTAGC 240
 TTGCATGCCT GCAGGTCGAC TTCCGCGCCC GCGGCGTCTG CCTTCGCCAG CAGGTTGTCC 300
 55 GCGGCCGCTG CCGGCCTGGT TCCGCGCCCG CCGCCTCGCG GCCAGCTCCC GCGCGGGCGC 360
 GTCCGCGTCC CCAACTCCGC GCGAAGACGG GCTCGTCCCA GAAGCGCAGC GGAAAGGCCG 420
 GCGTATAAAA TTTCGCTCGT CCGGTACAAA GACGCGGTCC GCGACTGCGT GGATGTCCAC 480
 60 GCCCAGGCAA GCAAACCTTA AACGCCCAGG CGCCATGGCC CCGATGCCGC CACAAAGAGC 540
 GCCGAAATTT CGCCCAGGCA CGCCGCGCCG CCCGACGCGT CTTTAGCGCA CCCGCCGGCG 600
 65 CTGTTGCCCC CGTGCTGCT GGCCGCCAC CGGCGGCCGC TGTCCCCGGC CTCAGCAGGG 660
 CCGGGGTCGC CGGCGGGCGG CCGCGGGGTG CGGCCACAGC CGCCCTTTTG CCCGTAGCCA 720

	GGGGAAGCGG CTGCCCCCTC TGCCGCCGCG GCCGCGGTTG CTCGGCTTTG CGTTTGCCCC	780
	GCGGCGATCG CCCCCTCGC CGCGAACGCG CGCGCGCGAA TGGGGCGTAC TCGGCGAGCC	840
5	CGGCTATTAT AGCCTCAAGG CGCGCCGCGT TGCTAGCGAT CGTCTGGGCC GGCAGGCGCG	900
	TCACTCTGAG CACGCGCATG CCCCCTGGG AGACGAACAC CTGCACCGGC GCTAGGACCA	960
	CCGGGTCTGG GCCCGGGGG GCGAGATCGC GCACAAGCCG GGCCGAGTCG CGCAGCTGCC	1020
10	GCAGCCCCC GAGGCGCTGG TCCATCTTGC TGGGCGTGT CATGTTTCGT GAAAAACGGC	1080
	ACGTCTTCAG CTCCACGATA AGACAGACGG CCCGGGCGTG CCCTGCCTCC GCGACCCGGA	1140
15	GTAGGCACAC GCAATCGGGC CGCCGGCTTT GCAGGTTTAC CTCAAAGCTC AGAGACACGC	1200
	CCACGACCTG CTTAAAAACC TCCGGGGCGC CAAACTTGCC CAAAAGCTGG GCGAGGCGCG	1260
	GGCGCAGCTT CTGCGCGCCA ACCGCCGCGC GTGCGTCGCA AGCCAGCGCC TCGTAAAAGC	1320
20	GGCTGTGGCA CCGGATCCCG GCGCGCAGGC GCGCACGTCG GTCGCGGTCG CGCGCCATGG	1380
	CCGAGCCCGC GCGCGCTCTC CGCGTCGTGC GTATCTACCT GGACGGCGCG CACGGGCAGG	1440
25	GAAAGACAAC AACGGGCCG GCGCTCGCG CCGCTTCCAC CGCTGGGGAG GGCCTGCTCT	1500
	TTTTCCCGGA GCCGATGGCG TACTGGCGCA CGATGTTTGG TACGGACGCC TTAAGTGGGA	1560
	TCCTCGCGGC GTCTGCGCGA TGCGCCGAG CCTCGCACGG GAGCGCACGC GCGCGCGGG	1620
30	CCGGCGCACC GCGCAGACGC GGACGCGGCG GGCCTGGTTG CGTACTACCA GGCCAGGTTT	1680
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35	CCGGGCGGCG AGCTGGTGGA CCCTCGTGTT CGACCGCCAC CCCGTGGCGC GCGTGCCTCT	1800
	GCTACCCCTT CGCCCGCTAC TGCCTCCGCG AGATCAACGC GGAAGATCTC AATTCTATGA	1860
40	TTTCTATCAT TACTTCCTCA CATGTTGGAG GCATTTTCTC TCCCTCTGCA CTTAATAGCC	1920
	TATCTTGCTT TAATTCTTC CCACTCTTGG AGGCTAGGTT TGGTTTGGTG GGCTGATGAG	1980
	GGAGGGAGAG ACCGCTCCAA GTACTTTAGC GGGTGGGATT GAAGCGGAGC CCTCCTGAGC	2040
45	TATGAGTGTC CTATGAGTGG GGCTGGAAC AAGAACCAGG GGCGTGGACA GGGTGTGTCA	2100
	CAGAGAAGGG GATGTGCCTG CTTCTTTCTG GCCCAGGAGG AACCAGGTCA ATTCTTCAGC	2160
	ACCTGGGTAC CCATAGAGCC CACCGCATCC CCAGCATGCC TGCTATTGTC TTCCAATCC	2220
50	TCCCCCTTGC TGTCTGCCC CACCCACCC CCCAGAATAG AATGACACCT ACTCAGACAA	2280
	TGCGATGCAA TTTCTCATT TTATTAGGAA AGGACAGTGG GAGTGGCACC TTCCAGGGTC	2340
55	AAGGAAGGCA CGGGGGAGGG GCAAACAACA GATGGCTGGC AACTAGAAGG CACAGCGGAT	2400
	CTGAGCTTGC ATGCCTGAGG TCGACCCTGG ATAAGCTGAT CCTCAATCAA TCAAGGTGGT	2460
	ATAAGAGTAA GACCACTTC TTTACAGCCT CCTCTCTTAG CAGTAGGTAT AACAACAAGA	2520
60	AATATGTCAC CACTTCAATA CTGTCATGTG TTAGCAAGTT ACCCATCATC ACCACTTCCC	2580
	CTGCCCCATA TTGGGTCCCC AAGGCCTTTT GTTCTGATAG GACCATGTAT GTTACCAGTA	2640
65	ACCAGAGCAC GTATCTTCCA CCCAGTAAAG CTACCACCAC CACCAATATG GACTCGGCGA	2700
	AGTAATCCCG ATGATGGTCA GTGACCTCCA GGTGGAACCA GTATTGATAA TCTCCTTTTA	2760

	GCATGTATTG CTGGAAGTAA CTGTCTCTGG GCTCAAAATA TTTATTTTTT AATGTCCTCG	2820
	TGTAGTTGAA GGTGCATGCC GTCTTTTCTA CAGGCCCTC ACTTGGTATG ATCTCATATG	2880
5	GCTTGCAAGG CATAGGCCCA AGTTTGGTGT CAAGAGCTAT GACCTGTACG ATTGTGTCTC	2940
	CTATCTTACA CTTTACCAAT CCTTGTGGTA CTATGGCTAC ACCCTCTCTG TTGCAAGTGG	3000
10	TGCCGTCTAC AAATCTGTAG CCAGTCTCAT TCTTCAACCT AACTTGCCG ATGGGGTAGT	3060
	GTGGCAACCC CTCACTTTTT TGAATTTAT AACCACACCA CTTGCAAGAT TCGACAGGGC	3120
	CTCTGTGTA TTGTAGCTGG TCCCAGTCA CACAAGTCCA ATTCCCTCCA AGATCACAGT	3180
15	TATAGAGATC CTCCCCCAGA GTTCTTTGGG TGATACAACC TTGTCTATAA GGAATGGTT	3240
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20	GTGTTGTATT GAACTTCCCT CTTACTACGG GATTGGCATC GCATGGGCAG AGTCCAAATT	3420
	CAAAGTCGTC AGCCATTTC AATGTTTCCC CTTGCCTTTG CCCCTCGAAA AGTTTTTTGA	3480
25	ATACCACACT GGTCTGCAAG GCTCTTGAAT GCAGAATTGC AAGATATCTA GTTCTCTTG	3540
	TGCACCTTGA GAGGTATGTA AACTTACCGT CTTTGACCA AGCTATGACC ATTGTGTCTT	3600
	CCAGCTTCAT TTCATGTGAG TAATCCTTCC AAACAGTGGT GAGGCCTTCA GCTCCTAATG	3660
30	GGCCAAATTCT ATCATTCCCTG GCTATGGCGT ATGAGTGTTT AGGTTTGCAG TCAATGTCCC	3720
	CTTGTAACCC TGTTATCAGT AGTAGCCATA GGATCCCCGA CGGCGCCGCG GCGATGGCCG	3780
35	CCGCGTAGAG CGCCAGCAGA GCGAGCATCG CACGCGCGAG CGAGGCCATG GTCGAAGCTT	3840
	GCCGCGGAGG CTGGATCGGT CCCGGTGTCT TCTATGGAGG TCAAAACAGC GTGGATGGCG	3900
40	TCTCCAGGCG ATCTGACGGT TACTAAACG AGCTCTGCTT ATATAGACCT CCCACCGTAC	3960
	ACGCCTACCG CCCATTGCG TCAATGGGGC GGAGTTGTTA CGACATTTTG GAAAGTCCCG	4020
	TTGATTTTGG TGCCAAAACA AACTCCCAT TACGTCAATG GGGTGGAGAC TTGGAAATCC	4080
45	CCGTGAGTCA AACCGCTATC CACGCCCAT TATGTACTGC CAAAACGCA TCACCATGGT	4140
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50	CTGGGCATAA TGCCAGGCGG GCCATTTACC GTCATTGACG TCAATAGGGG GCGTACTTGG	4260
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	ACGTCAATGG AAAGTCCCTA TTGGCGTTAC TATGGGAACA TACGTCATTA TTGACGTCAA	4380
55	TGGGCGGGGG TCGTTGGGCG GTCAGCCAGG CGGGCCATTT ACCGTAAGTT ATGTAACGCG	4440
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60	CAATAATCAA TGTCAACATG GCGGTAATGT TGGACATGAG CCAATATAAA TGTACATATT	4560
	ATGATATGGA TACAACGTAT GCAATGGCCA ATAGCCAATA TTGATTTATG CTATATAACC	4620
	AATGAATAAT ATGGCTAATG GCCAATATTG ATTCAATGTA TAGATCGATA TGCATTGGCC	4680
65	ATGTGCCAGC TTGATGTCGC CTCTATCGGC GATATAGCCT CATATCGTCT GTCACCTATA	4740
	TCGAAACTGC GATATTTGCG ACACACAGAA TCGCCCAAGT CACCAAAGGC GTCTATCGCC	4800

	ATCCCCCGTA AACGATATAA GCGTATCGCC AGATATCGCG TATGCCCAAA AATCAACTTT	4860
	TGGAAAAATG GCGATATCAG TTACACAGAA ACTCACATCG GCGACATTTT CAATATGCCA	4920
5	TATTTTCAAA TATCGATTTT TCCAATATCG CCATCTCTAT CGGCGATAAA CACCACTATC	4980
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10	TTTATTATTC ACTGCAGGTC GAGGAATTCG GATCTCGAGG TGTTCGTGCT GGACGTGTCC	5100
	GCGGCGCCAG ACGCGTGC GC GCGCGCCGTA CTGGACATGC GGCCCGCCAT GCAGGCCGCT	5160
	TGCGCGGACG GGGCGGCGGG GCGGACGCTG GCGACCCTGG CGCGTCAGTT CGCGCTAGAG	5220
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	ATTGGCAGCG GCTGTCTCG CGCTCGCCGC GGGCGCCCCC GCCCGCGCC GCGGCGGGGG	5460
	CGCCGAAGCC AGGGCAGCAC AGAGACGCCC GATACGAAAT CGAAGAGTGG GAAATGGTGG	5520
25	TCGGAGCCGG GCCGGCCGTG CACACGTTCA CCATCCGCTG CCTCGGGCCG CGGGGCATTG	5580
	AGCGCGTGGC CCACATTGCA AACCTCAGCC GGCTGCTGGA CGGGTACATA GCGGTCCACG	5640
30	TTGACGTTGC GCGCACCTCT GGCCTGCGGG ACGCCATGTT TTTCTGCCG CGCGCGGCCG	5700
	TCGACTCTAG AGGATCCCCG GGTACCGAGC TCGAATTCAC TGGCCGTCGT TTTACAACGT	5760
	CGTGACTGGG AAAACCCTGG CGTTACCCAA CTTAATCGCC TTGCAGCACA TCCCCCTTTC	5820
35	GCCAGCTGGC GTAATAGCGA AGAGGCCCGC ACCGATCGCC CTTCCCAACA GTTGCAGCAGC	5880
	CTGAATGGCG AATGGCGCCT GATGCGGTAT TTTCTCCTTA CGCATCTGTG CGGTATTTCA	5940
40	CACCGCATAT GGTGCACTCT CAGTACAATC TGCTCTGATG CCGCATAGTT AAGCCAGCCC	6000
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	TACAGACAAG CTGTGACCGT CTCCGGGAGC TGCATGTGTC AGAGGTTTTT ACCGTCATCA	6120
45	CCGAAACGCG CGAGACGAAA GGGCCTCGTG ATACGCCTAT TTTTATAGGT TAATGTCATG	6180
	ATAATAATGG TTTCTTAGAC GTCAGGTGGC ACTTTTCGGG GAAATGTGCG CGGAACCCCT	6240
50	ATTTGTTTAT TTTTCTAAAT ACATTCAAAT ATGTATCCGC TCATGAGACA ATAACCCTGA	6300
	TAAATGCTTC AATAATATTG AAAAAGGAAG AGTATGAGTA TTCAACATTT CCGTGTGCGC	6360
	CTTATTCCTT TTTTTCGCGC ATTTTGCCTT CCTGTTTTTG CTCACCCAGA AACGCTGGTG	6420
55	AAAGTAAAAG ATGCTGAAGA TCAGTTGGGT GCACGAGTGG GTTACATCGA ACTGGATCTC	6480
	AACAGCGGTA AGATCCTTGA GAGTTTTTCGC CCCGAAGAAC GTTTTCCAAT GATGAGCACT	6540
60	TTTAAAGTTC TGCTATGTGG CGCGGTATTA TCCCGTATTG ACGCCGGGCA AGAGCAACTC	6600
	GGTCGCCGCA TACACTATTC TCAGAATGAC TTGGTTGAGT ACTCACCAGT CACAGAAAAG	6660
	CATCTTACGG ATGGCATGAC AGTAAGAGAA TTATGCAGTG CTGCCATAAC CATGAGTGAT	6720
65	AACACTGCGG CCAACTTACT TCTGACAACG ATCGGAGGAC CGAAGGAGCT AACCGCTTTT	6780
	TTGCACAACA TGGGGGATCA TGTAACCTCGC CTTGATCGTT GGGAAACCGGA GCTGAATGAA	6840

GCCATACCAA ACGACGAGCG TGACACCACG ATGCCTGTAG CAATGGCAAC AACGTTGCGC 6900
 AACTATTAA CTGGCGAACT ACTTACTCTA GCTTCCCGGC AACAATTAAT AGACTGGATG 6960
 5 GAGGCGGATA AAGTTGCAGG ACCACTTCTG CGCTCGGCCC TTCCGGCTGG CTGGTTTATT 7020
 GCTGATAAAT CTGGAGCCGG TGAGCGTGGG TCTCGCGGTA TCATTGCAGC ACTGGGGCCA 7080
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 10 GAACGAAATA GACAGATCGC TGAGATAGGT GCCTCACTGA TTAAGCATTG GTAAGTGTCA 7200
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 15 ATCTAGGTGA AGATCCTTTT TGATAATCTC ATGACCAGAA TCCCTTAACG TGAGTTTTCG 7320
 TTCCACTGAG CGTCAGACCC CGTAGAAAAG ATCAAAGGAT CTTCTTGAGA TCCTTTTTTT 7380
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 20 CCGGATCAAG AGCTACCAAC TCTTTTTCCG AAGGTAACG GCTTCAGCAG AGCGCAGATA 7500
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 25 CCGCCTACAT ACCTCGCTCT GCTAATCCTG TTACCAGTGG CTGCTGCCAG TGGCGATAAG 7620
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 30 TACCTACAGC GTGAGCTATG AGAAAGCGCC ACGCTTCCCG AAGGGAGAAA GGCGGACAGG 7800
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 35 GCCTGGTATC TTTATAGTCC TGTCGGGTTT CGCCACCTCT GACTTGAGCG TCGATTTTGT 7920
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 40 TTCTGGCCT TTTGCTGGCC TTTTGCTCAC ATGTTCTTTC CTGCGTTATC CCCTGATTCT 8040
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 GAGCGCAGCG AGTCAGTGAG CGAGGAAGCG GAAGA 8135

45 (2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 8149 base pairs
 (B) TYPE: nucleic acid
 50 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

 (ii) MOLECULE TYPE: DNA (genomic)

 55 (iii) HYPOTHETICAL: NO
 (iv) ANTI-SENSE: NO

 (vi) ORIGINAL SOURCE:
 60 (A) ORGANISM: Bovine viral diarrhea virus
 (B) STRAIN: 2724
 (C) INDIVIDUAL ISOLATE: pBHvtkex-3::BGH/p53

 65 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

	GCGCCCAATA CGCAAACCGC CTCTCCCCGC GCGTTGGCCG ATTCATTAAT GCAGCTGGCA	60
	CGACAGGTTT CCCGACTGGA AAGCGGGCAG TGAGCGCAAC GCAATTAATG TGAGTTAGCT	120
5	CACTCATTAG GCACCCAGG CTTTACACTT TATGCTTCCG GCTCGTATGT TGTGTGGAAT	180
	TGTGAGCGGA TAACAATTTT ACACAGGAAA CAGCTATGAC CATGATTACG CCAAGCTAGC	240
10	TTGCATGCCT GCAGGTCGAC TTCCGCGCCC GCGGCGTCTG CCTTCGCCAG CAGGTTGTCC	300
	GCGGCCGCTG CCGGCCTGGT TCCGCGCCCC CCGCCTCGCG GCCAGCTCCC GCGCGGGCGC	360
	GTCCGCGTCC CCAACTCCGC GCGAAGACGG GCTCGTCCCA GAAGCGCAGC GGAAAGGCCG	420
15	GCGTATAAAA TTTCGCTCGT CCGGTACAAA GACGCGGTCC GCGACTGCGT GGATGTCCAC	480
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20	GCCGAAATTT CGCCCAGGCA CGCCGCGCCG CCCGACGCGT CTTTAGCGCA CCCGCCGGCG	600
	CTGTTGCCCC CGTGCTGCT GGCCGCCCAC CGGCGGCCGC TGTCCCCGGC CTCAGCAGGG	660
	CCGGGGTCGC CGGCGGGCGG CCGCGGGGTG CGGCCACAGC CGCCCTTTTG CCCGTAGCCA	720
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35	GCAGCCCCC GAGGCGCTGG TCCATCTTGC TGGGCGTGTT CATGTTCTGT GAAAAACGGC	1080
	ACGTCTTCAG CTCCACGATA AGACAGACGG CCCGGGCGTG CCCTGCCTCC GCGACCCGGA	1140
40	GTAGGCACAC GCAATCGGGC CGCCGGCTTT GCAGGTTTAC CTCAAAGCTC AGAGACACGC	1200
	CCACGACCTG CTTAAAAACC TCCGGGGCGC CAAACTTGCC CAAAAGCTGG GCGAGGCGCG	1260
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45	GGCTGTGGCA CCGGATCCCG GCGCGCAGGC GCGCACGTCG GTCGCGGTCG CGCGCCATGG	1380
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	TCCTCGCGGC GTCTGCGCGA TGCGCCGCAG CCTCGCACGG GAGCGCACGC GCGCGGCGGG	1620
55	CCGGCGCACC GCGCAGACGC GGACGCGGCG GGCCTGGTTG CGTACTACCA GGCCAGGTTT	1680
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65	ACTAAATTCA TGTCGCGCGA TAGTGGTGTT TATCGCCGAT AGAGATGGCG ATATTGGA	1980
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	GATATCGCCA TTTTCCAAA AGTTGATTTT TGGGCATACG CGATATCTGG CGATACGCTT	2100
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5	CAAATATCGC AGTTTCGATA TAGGTGACAG ACGATATGAG GCTATATCGC CGATAGAGGC	2220
	GACATCAAGC TGGCACATGG CCAATGCATA TCGATCTATA CATTGAATCA ATATTGGCCA	2280
10	TTAGCCATAT TATTCATTGG TTATATAGCA TAAATCAATA TTGGCTATTG GCCATTGCAT	2340
	ACGTTGTATC CATATCATAA TATGTACATT TATATTGGCT CATGTCCAAC ATTACCGCCA	2400
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15	AGCCCATATA TGGAGTTCCG CGTTACATAA CTTACGGTAA ATGGCCCGCC TGGCTGACCG	2520
	CCCAACGACC CCCGCCATT GACGTCAATA ATGACGTATG TTCCCATAGT AACGCCAATA	2580
	GGGACTTTCC ATTGACGTCA ATGGGTGGAG TATTTACGGT AACTGCCCCA CTGGCAGTA	2640
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	GCCTGGCATT ATGCCAGTA CATGACCTTA TGGGACTTTC CTAATTGGCA GTACATCTAC	2760
25	GTATTAGTCA TCGCTATTAC CATGGTGATG CGGTTTTGGC AGTACATCAA TGGGCGTGGA	2820
	TAGCGGTTTG ACTCACGGGG ATTTCCAAGT CTCCACCCCA TTGACGTCAA TGGGAGTTTG	2880
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30	CAAATGGGCG GTAGGCGTGT ACGGTGGGAG GTCTATATAA GCAGAGCTCG TTTAGTGAAC	3000
	CGTCAGATCG CCTGGAGACG CCATCCACGC TGTTTTGACC TCCATAGAAG ACACCGGGAC	3060
35	CGATCCAGCC TCCGCGGCAA GCTTCGACAT GATGGCTGCA GGCCCCGGA CCTCCCTGCT	3120
	CCTGGCTTTC GCCCTGCTCT GCCTGCCCTG GACTCAGGTG GTGGGCGCCT TCCCAGGGAT	3180
	CCTATGGCTA CTAATGATAA CAGGGGTACA AGGGGACATT GACTGCAAAC CTGAACACTC	3240
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45	AGACGGTAAG TTTACATACC TCTCAAGGTG CACAAGAGAA ACTAGATATC TTGCAATTCT	3420
	GCATTCAAGA GCCTTGACAG CCAGTGTGGT ATTCAAAAA CTTTTCGAGG GGCAAAGGCA	3480
	AGGGGAAACA TTTGAAATGG CTGACGACTT TGAATTTGGA CTCTGCCCCAT GCGATGCCAA	3540
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55	CACAGCAGTA GTGCGTGTGT ATAAGAGGTC CAAACCATTG CCTTATAGAC AAGGTTGTAT	3720
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	TGTGACTGGG GACCAGCTAC AATACACAGG AGGCCCTGTC GAATCTTGCA AGTGGTGTGG	3840
60	TTATAAATTC CAAAAAGTG AGGGGTGGCC ACACTACCCC ATCGGCAAGT GTAGGTTGAA	3900
	GAATGAGACT GGCTACAGAT TTGTAGACGG CACCACTTGC AACAGAGAGG GTGTAGCCAT	3960
65	AGTACCACAA GGATTGGTAA AGTGTAAGAT AGGAGACACA ATCGTACAGG TCATAGCTCT	4020
	TGACACCAAA CTTGGGCCTA TGCCTTGCAA GCCATATGAG ATCATACCAA GTGAGGGGCC	4080

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5	CCTGGAGGTC	ACTGACCATC	ATCGGGATTA	CTTCGCCGAG	TCCATATTGG	TGGTGGTGGT	4260
	AGCTTTACTG	GGTGAAGAT	ACGTGCTCTG	GTTACTGGTA	ACATACATGG	TCCTATCAGA	4320
	ACAAAAGGCC	TTGGGGACCC	AATATGGGGC	AGGGGAAGTG	GTGATGATGG	GTAACCTGCT	4380
10	AACACATGAC	AGTATTGAAG	TGGTGACATA	TTTCTTGTTG	TTATACCTAC	TGCTAAGAGA	4440
	GGAGGCTGTA	AAGAAGTGGG	TCTTACTCTT	ATACCACCTT	GATTGATTGA	GGATCAGCTT	4500
15	ATCCAGGGTC	GACCTCAGGC	ATGCAAGCTC	AGATCCGCTG	TGCCTTCTAG	TTGCCAGCCA	4560
	TCTGTTGTTT	GCCCCTCCCC	CGTGCCTTCC	TTGACCCCTG	AAGGTGCCAC	TCCCACCTGC	4620
	CTTTCCTAAT	AAAATGAGGA	AATTGCATCG	CATTGTCTGA	GTAGGTGTCA	TTCTATTCTG	4680
20	GGGGGTGGGG	TGGGGCAGGA	CAGCAAGGGG	GAGGATTGGG	AAGACAATAG	CAGGCATGCT	4740
	GGGGATGCGG	TGGGCTCTAT	GGGTACCCAG	GTGCTGAAGA	ATTGACCCGG	TTCCTCCTGG	4800
25	GCCAGAAAGA	AGCAGGCACA	TCCCCTTCTC	TGTGACACAC	CCTGTCCACG	CCCCTGGTTC	4860
	TTAGTTCCAG	CCCCACTCAT	AGGACACTCA	TAGCTCAGGA	GGGCTCCGCT	TCAATCCCAC	4920
	CCGCTAAAGT	ACTTGAGCG	GTCTCTCCCT	CCCTCATCAG	CCCACCAAAC	CAAACCTAGC	4980
30	CTCCAAGAGT	GGGAAGAAAT	TAAAGCAAGA	TAGGCTATTA	AGTGCAGAGG	GAGAGAAAAT	5040
	GCCTCCAACA	TGTGAGGAAG	TAATGATAGA	AATCATAGAA	TTGAGATCTC	GAGGTGTTCTG	5100
35	TGCTGGACGT	GTCCGCGGCG	CCAGACGCGT	GCGCGGCCGC	CGTACTGGAC	ATGCGGCCCCG	5160
	CCATGCAGGC	CGCTTGCGCG	GACGGGGCGG	CGGGCGCGAC	GCTGGCGACC	CTGGCGCGTC	5220
	AGTTCGCGCT	AGAGATGGCG	GGGGAGGCCA	CGGCGGGCCC	TAGGGGACTA	TAAAGCTGCC	5280
40	CCTGCGCTCG	CTCGCTCGCT	GCATTTGCGC	CCCGATCGCC	TTACGGGGAC	TCGGCGCTCG	5340
	GCGGATCCCC	TCCCGGCCCC	GCCGCGAAGC	AGGCCGCCAG	ACAAAAAAT	GCGGCGCCCCG	5400
45	CTCTGCGCGG	CGCTATTGGC	AGCGGCTGTC	CTCGCGCTCG	CCGCGGGCGC	CCCCGCCGCC	5460
	GCCCCGCGCG	GGGGCGCCGA	AGCCAGGGCA	GCACAGAGAC	GCCCCATACG	AAATCGAAGA	5520
	GTGGGAAATG	GTGGTCGGAG	CCGGGCCGGC	CGTGCACACG	TTCACCATCC	GCTGCCTCGG	5580
50	GCCGCGGGGC	ATTGAGCGCG	TGGCCACAT	TGCAAACCTC	AGCCGGCTGC	TGGACGGGTA	5640
	CATAGCGGTC	CACGTTGACG	TTGCGCGCAC	CTCTGGCCTG	CGGGACGCCA	TGTTTTTCCT	5700
55	GCCGCGCGCG	GCCGTCGACT	CTAGAGGATC	CCCGGTACC	GAGCTCGAAT	TCACTGGCCG	5760
	TCGTTTTTACA	ACGTCGTGAC	TGGGAAAACC	CTGGCGTTAC	CCAACTTAAT	CGCCTTGACG	5820
	CACATCCCCC	TTTCGCCAGC	TGGCGTAATA	GCGAAGAGGC	CCGCACCGAT	CGCCCTTCCC	5880
60	AACAGTTGCG	CAGCCTGAAT	GGCGAATGGC	GCCTGATGCG	GTATTTTCTC	CTTACGCATC	5940
	TGTGCGGTAT	TTCACACCGC	ATATGGTGCA	CTCTCAGTAC	AATCTGCTCT	GATGCCGCAT	6000
65	AGTTAAGCCA	GCCCCGACAC	CCGCCAACAC	CCGCTGACGC	GCCCTGACGG	GCTTGTCTGC	6060
	TCCCGGCATC	CGCTTACAGA	CAAGCTGTGA	CCGTCTCCGG	GAGCTGCATG	TGTCAGAGGT	6120

	TTTCACCGTC ATCACCGAAA CGCGCGAGAC GAAAGGGCCT CGTGATACGC CTATTTTTAT	6180
	AGGTTAATGT CATGATAATA ATGGTTTCTT AGACGTCAGG TGGCACTTTT CGGGGAAATG	6240
5	TGCGCGGAAC CCCTATTTGT TTATTTTTCT AAATACATTC AAATATGTAT CCGCTCATGA	6300
	GACAATAACC CTGATAAATG CTTCAATAAT ATTGAAAAAG GAAGAGTATG AGTATTCAAC	6360
10	ATTTCCGTGT CGCCCTTATT CCCTTTTTTG CGGCATTTTG CCTTCCTGTT TTTGCTCACC	6420
	CAGAAACGCT GGTGAAAGTA AAAGATGCTG AAGATCAGTT GGGTGCACGA GTGGGTTACA	6480
	TCGAACGGGA TCTCAACAGC GGTAAGATCC TTGAGAGTTT TCGCCCCGAA GAACGTTTTT	6540
15	CAATGATGAG CACTTTTAAA GTTCTGCTAT GTGGCGCGGT ATTATCCCGT ATTGACGCCG	6600
	GGCAAGAGCA ACTCGGTCGC CGCATACACT ATTCTCAGAA TGACTTGGTT GAGTACTCAC	6660
	CAGTCACAGA AAAGCATCTT ACGGATGGCA TGACAGTAAG AGAATTATGC AGTGCTGCCA	6720
20	TAACCATGAG TGATAACACT GCGGCCAACT TACTTCTGAC AACGATCGGA GGACCGAAGG	6780
	AGCTAACCGC TTTTTTGCAC AACATGGGGG ATCATGTAAC TCGCCTTGAT CGTTGGGAAC	6840
25	CGGAGCTGAA TGAAGCCATA CCAAACGACG AGCGTGACAC CACGATGCCT GTAGCAATGG	6900
	CAACAACGTT GCGCAAACTA TTAAGTGGCG AACTACTTAC TCTAGCTTCC CGGCAACAAT	6960
	TAATAGACTG GATGGAGGCG GATAAAGTTG CAGGACCACT TCTGCGCTCG GCCCTTCCGG	7020
30	CTGGCTGGTT TATGCTGAT AAATCTGGAG CCGGTGAGCG TGGGTCTCGC GGTATCATTG	7080
	CAGCACTGGG GCCAGATGGT AAGCCCTCCC GTATCGTAGT TATCTACACG ACGGGGAGTC	7140
35	AGGCAACTAT GGATGAACGA AATAGACAGA TCGTGAGAT AGGTGCCTCA CTGATTAAGC	7200
	ATTGGTAACT GTCAGACCAA GTTACTCAT ATATACTTTA GATTGATTTA AAACCTTCATT	7260
	TTTAATTTAA AAGGATCTAG GTGAAGATCC TTTTGTATAA TCTCATGACC AAAATCCCTT	7320
40	AACGTGAGTT TTCGTCCAC TGAGCGTCAG ACCCGTAGA AAAGATCAAA GGATCTTCTT	7380
	GAGATCCTTT TTTTCTGCGC GTAATCTGCT GCTTGCAAAC AAAAAACCA CCGCTACCAG	7440
45	CGGTGGTTTG TTTGCCGGAT CAAGAGCTAC CAACTCTTTT TCCGAAGGTA ACTGGCTTCA	7500
	GCAGAGCGCA GATACCAAAT ACTGTCCTTC TAGTGTAGCC GTAGTTAGGC CACCACTTCA	7560
	AGAACTCTGT AGCACCOCCT ACATACCTCG CTCTGCTAAT CCTGTTACCA GTGGCTGCTG	7620
50	CCAGTGGCGA TAAGTCGTGT CTTACCGGGT TGGACTCAAG ACGATAGTTA CCGGATAAGG	7680
	CGCAGCGGTC GGGCTGAACG GGGGGTTCGT GCACACAGCC CAGCTTGGAG CGAACGACCT	7740
55	ACACCGAACT GAGATACCTA CAGCGTGAGC TATGAGAAAG CGCCACGCTT CCCGAAGGGA	7800
	GAAAGGCGGA CAGGTATCCG GTAAGCGGCA GGGTCGGAAC AGGAGAGCGC ACGAGGGAGC	7860
	TTCCAGGGGG AAACGCCTGG TATCTTTATA GTCCTGTCCG GTTTCGCCAC CTCTGACTTG	7920
60	AGCGTCGATT TTTGTGATGC TCGTCAGGGG GCGGAGCCT ATGGAAAAAC GCCAGCAACG	7980
	CGGCCTTTTT ACGGTTCTTG GCCTTTTGCT GGCCTTTTGC TCACATGTTT TTTCTGCGT	8040
65	TATCCCCTGA TTCTGTGGAT AACCCTATTA CCGCCTTTGA GTGAGCTGAT ACCGCTCGCC	8100
	GCAGCCGAAC GACCGAGCGC AGCGAGTCAG TGAGCGAGGA AGCGGAAGA	8149

(2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 8135 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
- (v) ORIGINAL SOURCE:
 (A) ORGANISM: Bovine viral diarrhea virus
 (B) STRAIN: 2724
 (C) INDIVIDUAL ISOLATE: pBHVtkex-3::gIII/p53

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

	GCGCCCAATA CGCAAACCGC CTCTCCCCGC GCGTTGGCCG ATTCATTAAT GCAGCTGGCA	60
25	CGACAGGTTT CCCGACTGGA AAGCGGGCAG TGAGCGCAAC GCAATTAATG TGAGTTAGCT	120
	CACTCATTAG GCACCCCAGG CTTTACACTT TATGCTTCCG GCTCGTATGT TGTGTGGAAT	180
	TGTGAGCGGA TAACAATTTC ACACAGGAAA CAGCTATGAC CATGATTACG CCAAGCTAGC	240
30	TTGCATGCCT GCAGGTCGAC TTCCGCGCCC GCGGCGTCTG CCTTCGCCAG CAGGTTGTCC	300
	GCGGCCGCTG CCGGCCTGGT TCCGCGCCCG CCGCCTCGCG GCCAGCTCCC GCGCGGGCGC	360
35	GTCCGCGTCC CCAACTCCGC GCGAAGACGG GCTCGTCCCA GAAGCGCAGC GGAAAGGCCG	420
	GCGTATAAAA TTTCGCTCGT CCGGTACAAA GACGCGGTCC GCGACTGCGT GGATGTCCAC	480
	GCCCAGGCAA GCAAACTCTA AACGCCCCGAG CGCCATGGCC CCGATGCCGC CACAAAGAGC	540
40	GCCGAAATTT CGCCCAGGCA CGCCGCGCCG CCCGACGCGT CTTTAGCGCA CCCGCCGGCG	600
	CTGTTGCCCC CGTGCCCTGCT GGCCGCCCCAC CGGCGGCCGC TGTCCCCGGC CTCAGCAGGG	660
45	CCGGGGTCGC CGGCGGGCGG CCGCGGGGTG CGGCCACAGC CGCCCTTTTG CCCGTAGCCA	720
	GGGGAAGCGG CTGCCCTTTC TGCCGCCGCG GCCCGGGTTG CTCGGCTTTG CGTTTGCCCC	780
	GCGGCGATCG CCCCCTCGC CGCGAACGCG CGCGCGCGAA TGGGGCGTAC TCGGCGAGCC	840
50	CGGCTATTAT AGCCTCAAGG CGCGCCGCGT TGCTAGCGAT CGTCTGGGCC GGCAGGCGCG	900
	TCACTCTGAG CACGCGCATG CCCCCTGGG AGACGAACAC CTGCACCGGC GCTAGGACCA	960
55	CCGGGTCTGG GCCCGGGGGG GCGAGATCGC GCACAAGCCG GGCCGAGTCG CGCAGCTGCC	1020
	GCAGCCCCCC GAGGCGCTGG TCCATCTTGC TGGGCGTGTT CATGTTCTGT GAAAAACGGC	1080
	ACGTCTTCAG CTCCACGATA AGACAGACGG CCCGGGCGTG CCCTGCCTCC GCGACCCGGA	1140
60	GTAGGCACAC GCAATCGGGC CGCCGGCTTT GCAGGTTTAC CTCAAAGCTC AGAGACACGC	1200
	CCACGACCTG CTTAAAAACC TCCGGGGCGC CAAACTTGCC CAAAAGCTGG GCGAGGCGCG	1260
65	GGCGCAGCTT CTGCGCGCCA ACCGCCGCGC GTGCGTCGCA AGCCAGCGCC TCGTAAAGC	1320
	GGCTGTGGCA CCGGATCCCG GCGCGCAGGC GCGCACGTCG GTCGCGGTCT GCGCCCATGG	1380

	CCGAGCCCGC	GCGCGCTCTC	CGCGTCGTGC	GTATCTACCT	GGACGGCGCG	CACGGGCAGG	1440
	GAAAGACAAC	AACGGGGCCG	GCGCTCGCGG	CCGCTTCCAC	CGCTGGGGAG	GGCGTGCTCT	1500
5	TTTTCCCGGA	GCCGATGGCG	TACTGGCGCA	CGATGTTTGG	TACGGACGCC	TTAAGTGGGA	1560
	TCCTCGCGGC	GTCTGCGCGA	TGCGCCGCAG	CCTCGCACGG	GAGCGCACGC	GCGCGGCGGG	1620
	CCGGCGCACC	GCGCAGACGC	GGACGCGGCG	GGCCTGGTTG	CGTACTACCA	GGCCAGGTTC	1680
10	GCGGCCCCGT	ACTTAATTTT	GCACGCGCGT	GTCCGCGCTG	CTGCGCCGCC	TGGGCGGCGG	1740
	CCGGGCGGCG	AGCTGGTGGA	CCCTCGTGTT	CGACCGCCAC	CCCGTGGCGC	GCGTGCCCTCT	1800
15	GCTACCCCTT	CGCCCGCTAC	TGCCTCCGCG	AGATCAACGC	GGAAGATCCG	AATTCCTCGA	1860
	CCTGCAGTGA	ATAATAAAAT	GTGTGTTTGT	CCGAAATACG	CGTTTGAGAT	TTCTGTCCCG	1920
	ACTAAATTCA	TGTCGCGCGA	TAGTGGTGTT	TATCGCCGAT	AGAGATGGCG	ATATTGGAAA	1980
20	AATCGATATT	TGAAAATATG	GCATATTGAA	AATGTCGCCG	ATGTGAGTTT	CTGTGTA ACT	2040
	GATATCGCCA	TTTTTCCAAA	AGTTGATTTT	TGGGCATACG	CGATATCTGG	CGATACGCTT	2100
25	ATATCGTTTA	CGGGGGATGG	CGATAGACGC	CTTTGGTGAC	TTGGGCGATT	CTGTGTGTCG	2160
	CAAATATCGC	AGTTTCGATA	TAGGTGACAG	ACGATATGAG	GCTATATCGC	CGATAGAGGC	2220
	GACATCAAGC	TGGCACATGG	CCAATGCATA	TCGATCTATA	CATTGAATCA	ATATTGGCCA	2280
30	TTAGCCATAT	TATTCATTGG	TTATATAGCA	TAAATCAATA	TTGGCTATTG	GCCATTGCAT	2340
	ACGTTGTATC	CATATCATAA	TATGTACATT	TATATTGGCT	CATGTCCAAC	ATTACCGCCA	2400
35	TGTTGACATT	GATTATTGAC	TAGTTATTAA	TAGTAATCAA	TTACGGGGTC	ATTAGTTTCAT	2460
	AGCCCATATA	TGGAGTTCCG	CGTTACATAA	CTTACGGTAA	ATGGCCCGCC	TGGCTGACCG	2520
	CCCAACGACC	CCCGCCATT	GACGTCAATA	ATGACGTATG	TTCCCATAGT	AACGCCAATA	2580
40	GGGACTTTCC	ATTGACGTCA	ATGGGTGGAG	TATTTACGGT	AACTGCCCCA	CTTGGCAGTA	2640
	CATCAAGTGT	ATCATATGCC	AAGTACGCCC	CCTATTGACG	TCAATGACGG	TAAATGGCCC	2700
45	GCCTGGCATT	ATGCCCAGTA	CATGACCTTA	TGGGACTTTC	CTACTTGGCA	GTACATCTAC	2760
	GTATTAGTCA	TCGCTATTAC	CATGGTGATG	CGGTTTGGC	AGTACATCAA	TGGGCGTGGA	2820
	TAGCGGTTTG	ACTCACGGGG	ATTTCCAAGT	CTCCACCCCA	TTGACGTCAA	TGGGAGTTTG	2880
50	TTTTGGCACC	AAAATCAACG	GGACTTTCCA	AAATGTCGTA	ACAACTCCGC	CCCATTGACG	2940
	CAAATGGGCG	GTAGGCGTGT	ACGGTGGGAG	GTCTATATAA	GCAGAGCTCG	TTTAGTGAAC	3000
55	CGTCAGATCG	CCTGGAGACG	CCATCCACGC	TGTTTTGACC	TCCATAGAAG	ACACCGGGAC	3060
	CGATCCAGCC	TCCGCGGCAA	GCTTCGACCA	TGGCCTCGCT	CGCGCGTGCG	ATGCTCGCTC	3120
	TGCTGGCGCT	CTACGCGGCG	GCCATCGCCG	CGGCGCCGTC	GGGGATCCTA	TGGCTACTAC	3180
60	TGATAACAGG	GGTACAAGGG	GACATTGACT	GCAAACCTGA	ACACTCATAC	GCCATAGCCA	3240
	GGAATGATAG	AATTGGCCCA	TTAGGAGCTG	AAGGCCTCAC	CACTGTTTGG	AAGGATTACT	3300
65	CACATGAAAT	GAAGCTGGAA	GACACAATGG	TCATAGCTTG	GTGCAAAGAC	GGTAAGTTTA	3360
	CATACCTCTC	AAGGTGCACA	AGAGAAACTA	GATATCTTGC	AATTCTGCAT	TCAAGAGCCT	3420

	TGCAGACCAG	TGTGGTATTC	AAAAAACTTT	TCGAGGGGCA	AAGGCAAGGG	GAAACATTTG	3480
	AAATGGCTGA	CGACTTTGAA	TTTGGACTCT	GCCCATGCGA	TGCCAATCCC	GTAAGTAAGAG	3540
5	GGAGTTCAA	TACAACACTG	CTAAACGGAC	CGGCCTTCCA	GATGGTATGC	CCTATAGGAT	3600
	GGACAGGAAC	TGTGAGCTGT	ATGTTAGCTA	ATAGGGACAC	CCTAGACACA	GCAGTAGTGC	3660
	GTGTGTATAA	GAGGTCCAAA	CCATTCCCTT	ATAGACAAGG	TTGTATCACC	CAAAGAACTC	3720
10	TGGGGGAGGA	TCTCTATAAC	TGTGATCTTG	GAGGGAATTG	GACTTGTGTG	ACTGGGGACC	3780
	AGCTACAATA	CACAGGAGGC	CCTGTCTGAAT	CTTGCAAGTG	GTGTGGTTAT	AAATTCCAAA	3840
15	AAAGTGAGGG	GTTGCCACAC	TACCCCATCG	GCAAGTGTAG	GTTGAAGAAT	GAGACTGGCT	3900
	ACAGATTTGT	AGACGGCACC	ACTTGCAACA	GAGAGGGTGT	AGCCATAGTA	CCACAAGGAT	3960
	TGGTAAAGTG	TAAGATAGGA	GACACAATCG	TACAGGTCAT	AGCTCTTGAC	ACCAAACCTG	4020
20	GGCCTATGCC	TTGCAAGCCA	TATGAGATCA	TACCAAGTGA	GGGGCCTGTA	GAAAGACGG	4080
	CATGCACCTT	CAACTACACG	AGGACATTAA	AAAATAAATA	TTTGAGCCC	AGAGACAGTT	4140
25	ACTTCCAGCA	ATACATGCTA	AAAGGAGATT	ATCAATACTG	GTTGACCTG	GAGGTCACTG	4200
	ACCATCATCG	GGATTACTTC	GCCGAGTCCA	TATTGGTGGT	GGTGGTAGCT	TTACTGGGTG	4260
	GAAGATACGT	GCTCTGGTTA	CTGGTAACAT	ACATGGTCCT	ATCAGAACAA	AAGGCCTTGG	4320
30	GGACCCAATA	TGGGGCAGGG	GAAGTGGTGA	TGATGGGTAA	CTTGCTAACA	CATGACAGTA	4380
	TTGAAGTGGT	GACATATTTT	TTGTTGTTAT	ACCTACTGCT	AAGAGAGGAG	GCTGTAAAGA	4440
35	AGTGGGTCTT	ACTCTTATAC	CACCTTGATT	GATTGAGGAT	CAGCTTATCC	AGGGTCGACC	4500
	TCAGGCATGC	AAGCTCAGAT	CCGCTGTGCC	TTCTAGTTGC	CAGCCATCTG	TTGTTTGCCC	4560
	CTCCCCCGTG	CCTTCCTTGA	CCCTGGAAGG	TGCCACTCCC	ACTGTCCCTT	CCTAATAAAA	4620
40	TGAGGAAATT	GCATCGCATT	GTCTGAGTAG	GTGTCATTCT	ATTCTGGGGG	GTGGGGTGGG	4680
	GCAGGACAGC	AAGGGGGAGG	ATTGGGAAGA	CAATAGCAGG	CATGCTGGGG	ATGCGGTGGG	4740
45	CTCTATGGGT	ACCCAGGTGC	TGAAGAATTG	ACCCGGTTCC	TCCTGGGCCA	GAAAGAAGCA	4800
	GGCACATCCC	CTTCTCTGTG	ACACACCCTG	TCCACGCCCC	TGGTTCTTAG	TTCCAGCCCC	4860
	ACTCATAGGA	CACTCATAGC	TCAGGAGGGC	TCCGCTTCAA	TCCCACCCGC	TAAAGTACTT	4920
50	GGAGCGGTCT	CTCCCTCCCT	CATCAGCCCA	CCAAACCAAA	CCTAGCCTCC	AAGAGTGGGA	4980
	AGAAATTAAA	GCAAGATAGG	CTATTAAGTG	CAGAGGGAGA	GAAAATGCCT	CCAACATGTG	5040
55	AGGAAGTAAT	GATAGAAATC	ATAGAATTGA	GATCTCGAGG	TGTTCTGTGCT	GGACGTGTCC	5100
	GCGGCGCCAG	ACGCGTGCGC	GGCCGCCGTA	CTGGACATGC	GGCCCGCCAT	GCAGGCCGCT	5160
	TGCGCGGACG	GGGCGGCGGG	CGCGACGCTG	GCGACCCTGG	CGCGTCAGTT	CGCGCTAGAG	5220
60	ATGGCGGGGG	AGGCCACGGC	GGGCCCTAGG	GGACTATAAA	GCTGCCCCCTG	CGCTCGCTCG	5280
	CTCGCTGCAT	TTGCGCCCCG	ATCGCCTTAC	GGGGACTCGG	CGCTCGGCGG	ATCCCCCTCC	5340
65	GGCCCCGCCG	CGAAGCAGGC	CGCCAGACAA	AAAAATGCGG	CGCCCGCTCT	GCGCGGCGCT	5400
	ATTGGCAGCG	GCTGTCCTCG	CGCTCGCCGC	GGGCGCCCCC	GCCGCGCCCC	GCGGCGGGGG	5460

	CGCCGAAGCC	AGGGCAGCAC	AGAGACGCCC	GATACGAAAT	CGAAGAGTGG	GAAATGGTGG	5520
	TCGGAGCCGG	GCCGGCCGTG	CACACGTTCA	CCATCCGCTG	CCTCGGGCCG	CGGGGCATTG	5580
5	AGCGCGTGGC	CCACATTGCA	AACCTCAGCC	GGCTGCTGGA	CGGGTACATA	GCGGTCCACG	5640
	TTGACGTTGC	GCGCACCTCT	GGCCTGCGGG	ACGCCATGTT	TTTCCTGCCG	CGCGCGGGCCG	5700
	TCGACTCTAG	AGGATCCCCG	GTACCGAGC	TCGAATTCAC	TGGCCGTCGT	TTTACAACGT	5760
10	CGTGACTGGG	AAAACCCTGG	CGTTACCCAA	CTTAATCGCC	TTGCAGCACA	TCCCCCTTTC	5820
	GCCAGCTGGC	GTAATAGCGA	AGAGGCCCCG	ACCGATCGCC	CTTCCCAACA	GTTGCGCAGC	5880
15	CTGAATGGCG	AATGGCGCCT	GATGCGGTAT	TTTCTCCTTA	CGCATCTGTG	CGGTATTTCA	5940
	CACCGCATAT	GGTGCACTCT	CAGTACAATC	TGCTCTGATG	CCGCATAGTT	AAGCCAGCCC	6000
	CGACACCCGC	CAACACCCGC	TGACGCGCCC	TGACGGGCTT	GTCTGCTCCC	GGCATCCGCT	6060
20	TACAGACAAG	CTGTGACCGT	CTCCGGGAGC	TGCATGTGTC	AGAGGTTTTT	ACCGTCATCA	6120
	CCGAAACGCG	CGAGACGAAA	GGGCCTCGTG	ATACGCCTAT	TTTTATAGGT	TAATGTCATG	6180
25	ATAATAATGG	TTTCTTAGAC	GTCAGGTGGC	ACTTTTTCGG	GAAATGTGCG	CGGAACCCCT	6240
	ATTTGTTTAT	TTTTCTAAAT	ACATTCAAAT	ATGTATCCGC	TCATGAGACA	ATAACCCTGA	6300
	TAAATGCTTC	AATAATATTG	AAAAAGGAAG	AGTATGAGTA	TTCAACATTT	CCGTGTCGCC	6360
30	CTTATTCCTT	TTTTTGCGGC	ATTTTGCTT	CCTGTTTTTG	CTCACCCAGA	AACGCTGGTG	6420
	AAAGTAAAAG	ATGCTGAAGA	TCAGTTGGGT	GCACGAGTGG	GTTACATCGA	ACTGGATCTC	6480
35	AACAGCGGTA	AGATCCTTGA	GAGTTTTTCG	CCCGAAGAAC	GTTTTCCAAT	GATGAGCACT	6540
	TTTAAAGTTC	TGCTATGTGG	CGCGGTATTA	TCCCGTATTG	ACGCCGGGCA	AGAGCAACTC	6600
	GGTCGCCGCA	TACACTATTC	TCAGAATGAC	TTGGTTGAGT	ACTCACCAGT	CACAGAAAAG	6660
40	CATCTTACGG	ATGGCATGAC	AGTAAGAGAA	TTATGCAGTG	CTGCCATAAC	CATGAGTGAT	6720
	AACACTGCGG	CCAACCTTACT	TCTGACAACG	ATCGGAGGAC	CGAAGGAGCT	AACCGCTTTT	6780
45	TTGCACAACA	TGGGGGATCA	TGTAACTCGC	CTTGATCGTT	GGGAACCGGA	GCTGAATGAA	6840
	GCCATACCAA	ACGACGAGCG	TGACACCACG	ATGCCTGTAG	CAATGGCAAC	AACGTTGCGC	6900
	AAACTATTAA	CTGGCGAACT	ACTTACTCTA	GCTTCCCGGC	AACAATTAAT	AGACTGGATG	6960
50	GAGGCGGATA	AAGTTGCAGG	ACCACTTCTG	CGCTCGGCCC	TTCCGGCTGG	CTGGTTTATT	7020
	GCTGATAAAT	CTGGAGCCGG	TGAGCGTGGG	TCTCGCGGTA	TCATTGCAGC	ACTGGGGCCA	7080
55	GATGGTAAGC	CCTCCCGTAT	CGTAGTTATC	TACACGACGG	GGAGTCAGGC	AACATATGGAT	7140
	GAACGAAATA	GACAGATCGC	TGAGATAGGT	GCCTCACTGA	TTAAGCATTG	GTAACGTGCA	7200
	GACCAAGTTT	ACTCATATAT	ACTTTAGATT	GATTTAAAC	TTCATTTTTA	ATTTAAAAGG	7260
60	ATCTAGGTGA	AGATCCTTTT	TGATAATCTC	ATGACCAAAA	TCCCTTAACG	TGAGTTTTCG	7320
	TTCCACTGAG	CGTCAGACCC	CGTAGAAAAG	ATCAAAGGAT	CTTCTTGAGA	TCCTTTTTTT	7380
65	CTGCGCGTAA	TCTGCTGCTT	GCAAACAAAA	AAACCACCGC	TACCAGCGGT	GGTTTGTGTTG	7440
	CCGGATCAAG	AGCTACCAAC	TCTTTTTCCG	AAGGTAAGTG	GCTTCAGCAG	AGCGCAGATA	7500

	CCAAATACTG TCCTTCTAGT GTAGCCGTAG TTAGGCCACC ACTTCAAGAA CTCTGTAGCA	7560
	CCGCCTACAT ACCTCGCTCT GCTAATCCTG TTACCAGTGG CTGCTGCCAG TGGCGATAAG	7620
5	TCGTGTCTTA CCGGGTTGGA CTCAAGACGA TAGTTACCGG ATAAGGCGCA GCGGTCGGGC	7680
	TGAACGGGGG GTTCGTGCAC ACAGCCCAGC TTGGAGCGAA CGACCTACAC CGAACTGAGA	7740
	TACCTACAGC GTGAGCTATG AGAAAGCGCC ACGCTTCCCG AAGGGAGAAA GGCGGACAGG	7800
10	TATCCGGTAA GCGGCAGGGT CGGAACAGGA GAGCGCACGA GGGAGCTTCC AGGGGGAAAC	7860
	GCCTGGTATC TTTATAGTCC TGTCGGGTTT CGCCACCTCT GACTTGAGCG TCGATTTTTG	7920
15	TGATGCTCGT CAGGGGGGCG GAGCCTATGG AAAACGCCA GCAACGCGGC CTTTTTACGG	7980
	TTCTTGGCCT TTTGCTGGCC TTTTGCTCAC ATGTTCTTTC CTGCGTTATC CCCTGATTCT	8040
	GTGGATAACC GTATTACCGC CTTTGAGTGA GCTGATACCG CTCGCCGAG CCGAACGACC	8100
20	GAGCGCAGCG AGTCAGTGAG CGAGGAAGCG GAAGA	8135

Claims

1. A replicating nonpathogenic virus, for preventing disease caused by Bovine Viral Diarrhea Virus (BVDV), where said replicating nonpathogenic virus comprises:
a gene or gene combination taken from a BVDV virus, and said replicating
5 nonpathogenic virus functionally expresses said gene or gene combination.
2. A virus of claim 1, where said replicating nonpathogenic virus is attenuated.
3. A virus of claim 2, where said replicating nonpathogenic virus is selected
10 from attenuated Bovine Herpes Virus type 1 (BHV-1), attenuated adenoviruses, attenuated bovine mammillitis virus, attenuated bovine papillomavirus, or attenuated pseudorabies virus.
4. A virus of claim 2, where said replicating nonpathogenic virus is attenuated
15 and contains and expresses any combination of the following genes: the genes that code for gp48, gp25, p14 capsid protein, p20 N-terminal protease and p125/p80 protein.
5. A virus of claim 3, where said replicating nonpathogenic virus is attenuated
20 and contains and expresses any combination of the following genes: the genes that code for gp48, gp25, p14 capsid protein, p20 N-terminal protease and p125/p80 protein.
6. A virus of claim 2, where said attenuation is created by making the
25 thymidine kinase (tk) gene nonfunctional.
7. A virus of claim 3, where said attenuation is created by making the thymidine kinase (tk) gene nonfunctional.
8. A virus of claim 4, where said attenuation is created by making the
30 thymidine kinase (tk) gene nonfunctional.
9. A virus of claim 5, where said attenuation is created by making the thymidine kinase (tk) gene nonfunctional.

10. A virus of claim 9, where said replicating nonpathogenic virus is attenuated Bovine Herpes Virus type 1 (BHV-1).
11. A virus of claim 10, where said replicating nonpathogenic virus contains and
5 expresses the gene that codes for gp53, a glycoprotein of the Bovine Viral Diarrhea Virus (BVDV).
12. A virus of claim 11, where a signal peptide is inserted preceeding the gene or gene combination that codes for gp53 in said Bovine Herpes Virus type 1 (BHV-1).
- 10 13. A virus of claim 12, where said gene that codes for gp53 is inserted into the inactivated thymidine kinase (tk) gene site.
14. A virus of claim 13, where the functionally expressing gene or gene
15 combination, used to create the virus, comprises a recombined plasmid with intact viral DNA, said plasmid comprising:
- a) a BHV-1 genomic DNA fragment containing the thymidine kinase (tk) gene and having a deletion to the thymidine kinase (tk) gene,
 - b) a promoter/polyadenylation signal inserted in the thymidine kinase (tk)
20 gene deletion,
 - c) a signal peptide gene sequence preceding a gp53 gene or gene combination all of which is inserted between the promoter and the polyadenylation signal.
15. A virus of claim 14, where said signal peptide gene sequence is taken from
25 any well characterized signal peptide sequences such as any of the thirty-nine examples of well characterized signal peptide sequences found in Perlman, D., et al., *J. Mol. Biol.* Vol. 167 pp. 391-409 (1983).
16. A virus of claim 15, where said signal peptide gene sequence is taken from
30 Psuedorabies Virus gIII gene (PRV) and/or Bovine Growth Hormone (BGH).
17. A virus of claim 16 where the plasmid is selected from the following plasmids,
- a) pBHVtkex-1::BGH/p53;
 - b) pBHVtkex-1::gIII/p53;
- 35

- c) pBHVtkex-3::BGH/p53; or
- d) pBHVtkex-3::gIII/p53.

18. A virus of claim 17, where said virus that produces the product of a
5 functionally expressing gene or gene combination is selected from one of the
following viruses,

T11-3, T11-6, or T11-8.

19. A virus of claim 18, where said gene or gene combination is T11-6.

10

20. A virus of claim 11, where said gene that codes for gp53 is inserted into the
inactivated thymidine kinase (tk) gene site.

21. A virus of claim 20, where the functionally expressing gene or gene
15 combination, used to create the virus, comprises a recombined plasmid with intact
viral DNA, said plasmid comprising:

a) a BHV-1 genomic DNA fragment containing the thymidine kinase (tk)
gene and having a deletion to the thymidine kinase (tk) gene,

b) a promoter/polyadenylation signal inserted in the thymidine kinase (tk)
20 gene deletion,

c) a gp53 gene or gene combination inserted between the promoter and the
polyadenylation signal.

22. A virus of claim 21, where said plasmid is made from a plasmid having the
25 characteristics of plasmid pHAS4.

23. A virus of claim 22, where said plasmid is pBHVtkex-3::p53.

24. A virus of claim 23, where said virus is selected from one of the following
30 viruses,

T2-3#3 or T2-2#5.

25. A vaccine for preventing disease caused by Bovine Viral Diarrhea Virus
(BDVD) comprising a pharmaceutically effective amount of a virus of claim 1 and a
35 carrier.

26. A vaccine as claimed in claim 25, for preventing disease caused by Bovine Viral Diarrhea Virus (BDVD) comprising a pharmaceutically effective amount of a virus of claim 1 and a carrier, said carrier comprising any physiological buffered medium, i.e. about pH 7.0 to 7.4 containing from about 2.5 to 15% serum which does
5 not contain antibodies to BHV.

27. A method of immunizing an animal against infectious disease caused by Bovine Viral Diarrhea Virus (BDVD) comprising administering to an animal a pharmaceutically effective amount of a virus of claim 1.

10

28. A process of preparing a virus of claim 1 comprising:
a) isolation of a functionally expressing gene or gene combination that causes BVDV,
b) inserting said gene or gene combination into a replicating nonpathogenic
15 virus,
c) selecting a live-virus that functionally expresses the product of said gene or gene combination.

29. A method of preparing a virus of claim 11 where the functionally expressing
20 gene or gene combination, used to create the virus, is produced by a process comprising the recombination of a plasmid with intact viral DNA, said plasmid comprising:

a) a BHV-1 genomic DNA fragment containing the thymidine kinase (tk) gene and having a deletion to the thymidine kinase (tk) gene,
25 b) inserting into the thymidine kinase (tk) gene deletion of said plasmid a promoter/polyadenylation signal,
c) inserting a gp53 gene or gene combination between the promoter and the polyadenylation signal,
d) transfecting cells with said plasmid to produce a recombinant virus
30 containing said functional gene or gene combination inserted into a live virus that does not cause immunosuppression in the usual host and expressing said functional gene or gene combination.

30. A method of preparing a virus of claim 12 where the functionally expressing
35 gene or gene combination, used to create the virus, is produced by a process

comprising the recombination of a plasmid with intact viral DNA, said plasmid comprising:

- a) a BHV-1 genomic DNA fragment containing the thymidine kinase (tk) gene and having a deletion to the thymidine kinase (tk) gene,
- 5 b) inserting into the thymidine kinase (tk) gene deletion of said plasmid a promoter/polyadenylation signal,
- c) inserting a gp53 gene or gene combination preceded by a signal peptide gene sequence between the promoter and the polyadenylation signal,
- d) transfecting cells with said plasmid to produce a recombinant virus
- 10 containing said functional gene or gene combination inserted into a live virus that does not cause immunosuppression in the usual host and expressing said functional gene or gene combination.

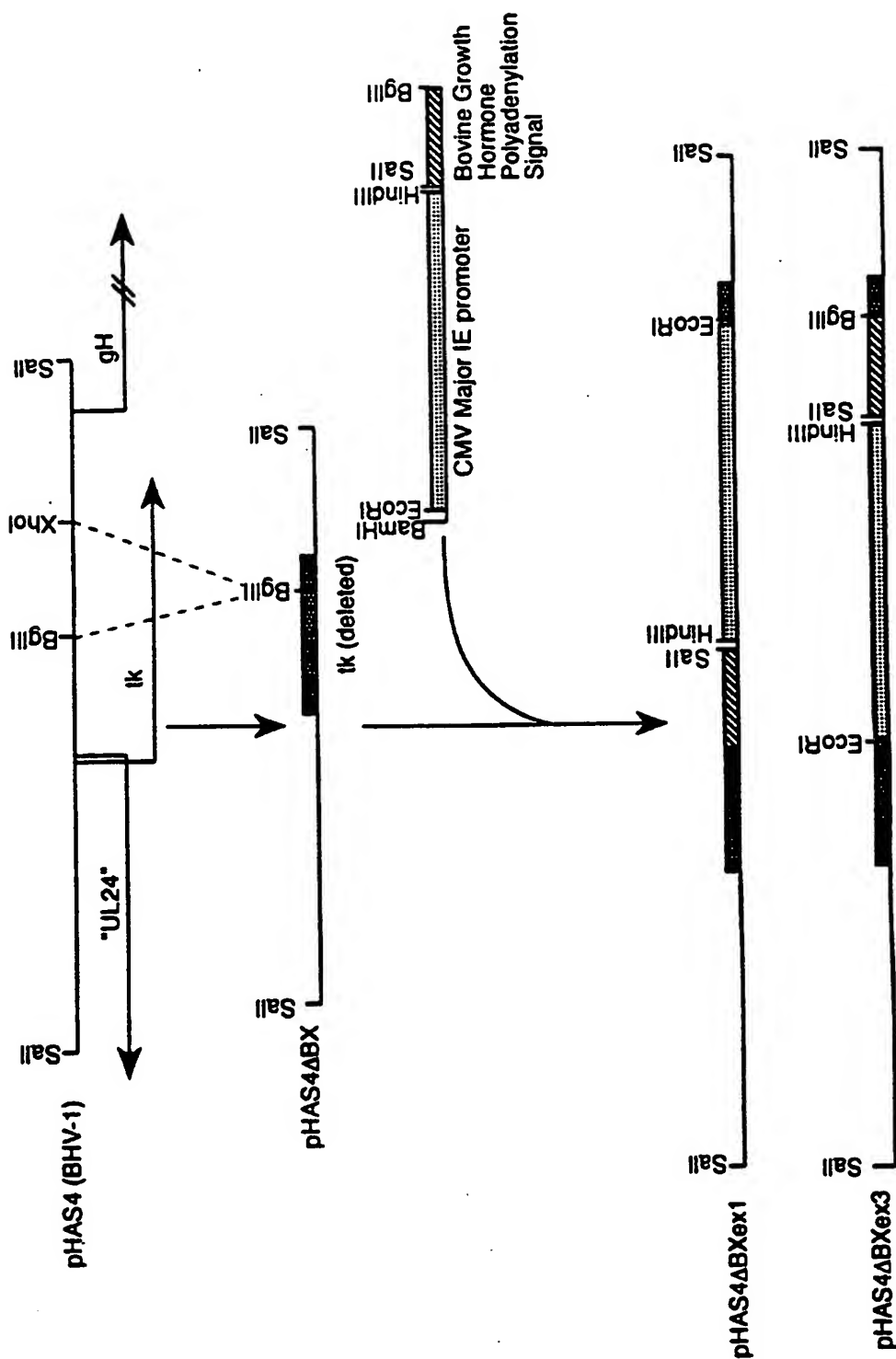
FIGURE 1

FIGURE 2A

BGH

Sali

```
TCGACATGATGGCTGCAGGCCCGGACCTCCCTGCTCCTGGCTTTCCGCCCTGCTCTGC
-----+-----+-----+-----+-----+-----+-----+
GTACTACCGACGTCCGGGGGCTGGAGGGACGAGGACCGAAAGCGGGACGAGACG
M M A A G P T Y S L L L A F A L L C
```

BamHI

```
CTGCCCTGGACTCAGGTGGTGGGCGCCTTCCCAGGG
-----+-----+-----+-----+-----+
GACGGGACCTGAGTCCACCACCCGCGGAAGGGTCCCTAG
L P W T Q V V G A F P G
```

PRV gIII

Sali

```
TCGACCATGGCCTCGCTCGCGCGTGCGATGCTCGCTCTGCTGGCGCTCTACGGGCGGC
-----+-----+-----+-----+-----+-----+
GGTACCGGAGCGAGCGCGCACGCTACGAGCGAGACGACCGCGAGATGCGCCGCCG
M A S L A R A M L A L L A L Y A A A
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BamHI

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CATCGCCGCGGCGCCGTCGGGG
-----+-----+-----+
GTAGCGGCGCCGCGGCAGCCCCTAG
I A A A P S G
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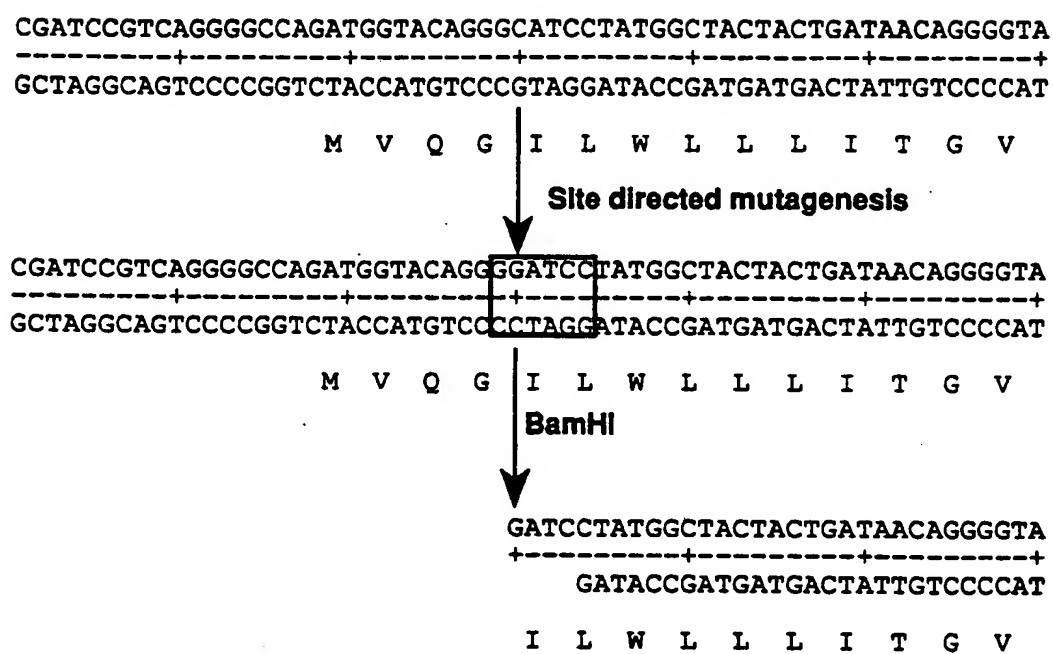
FIGURE 2B

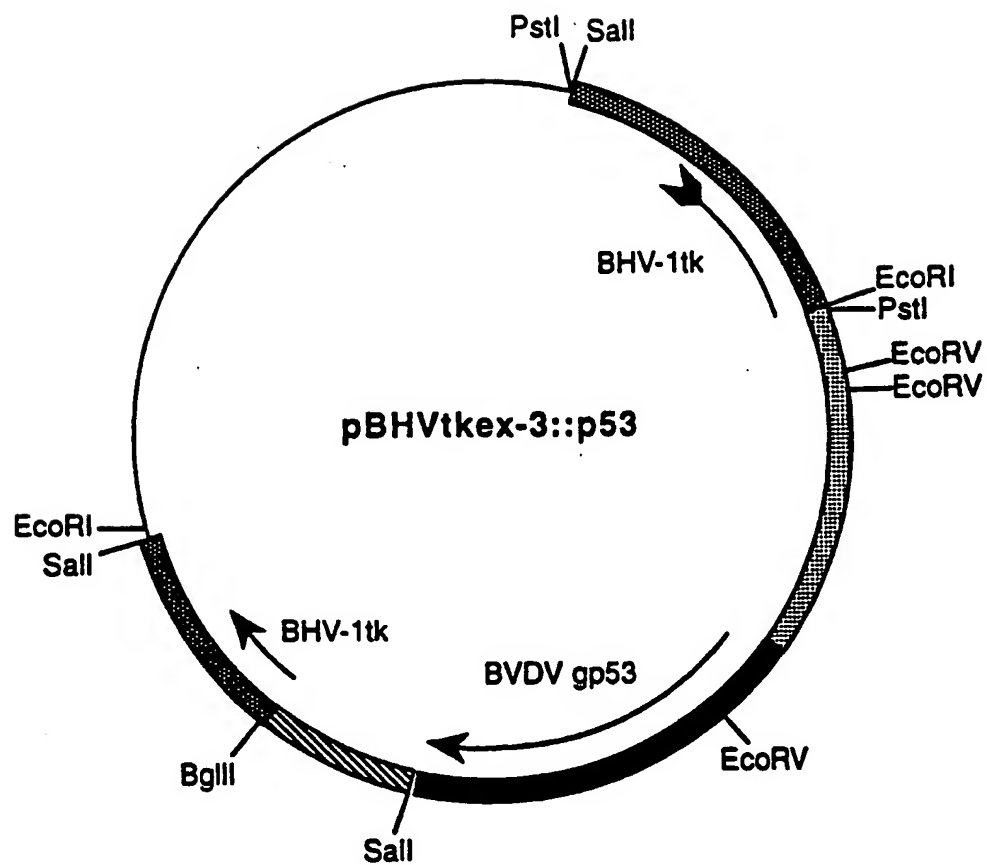
FIGURE 3A

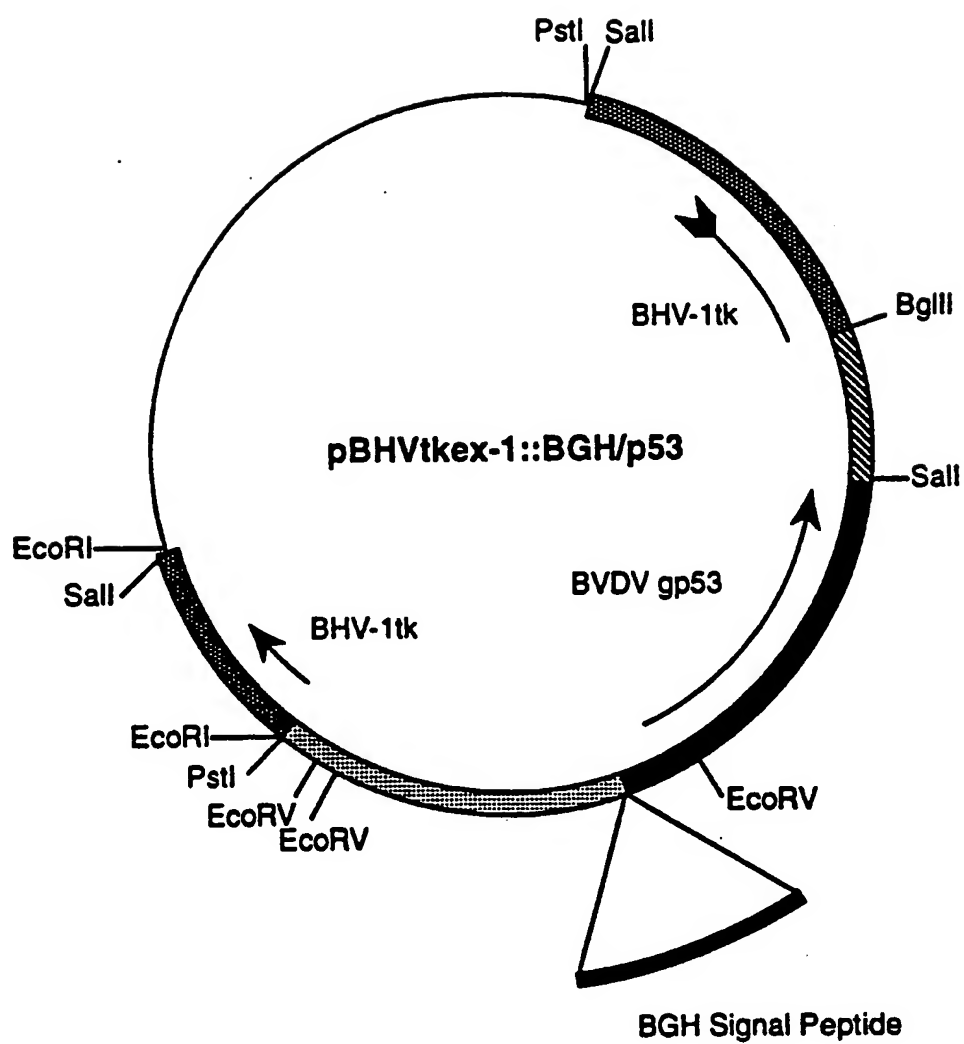
FIGURE 3B

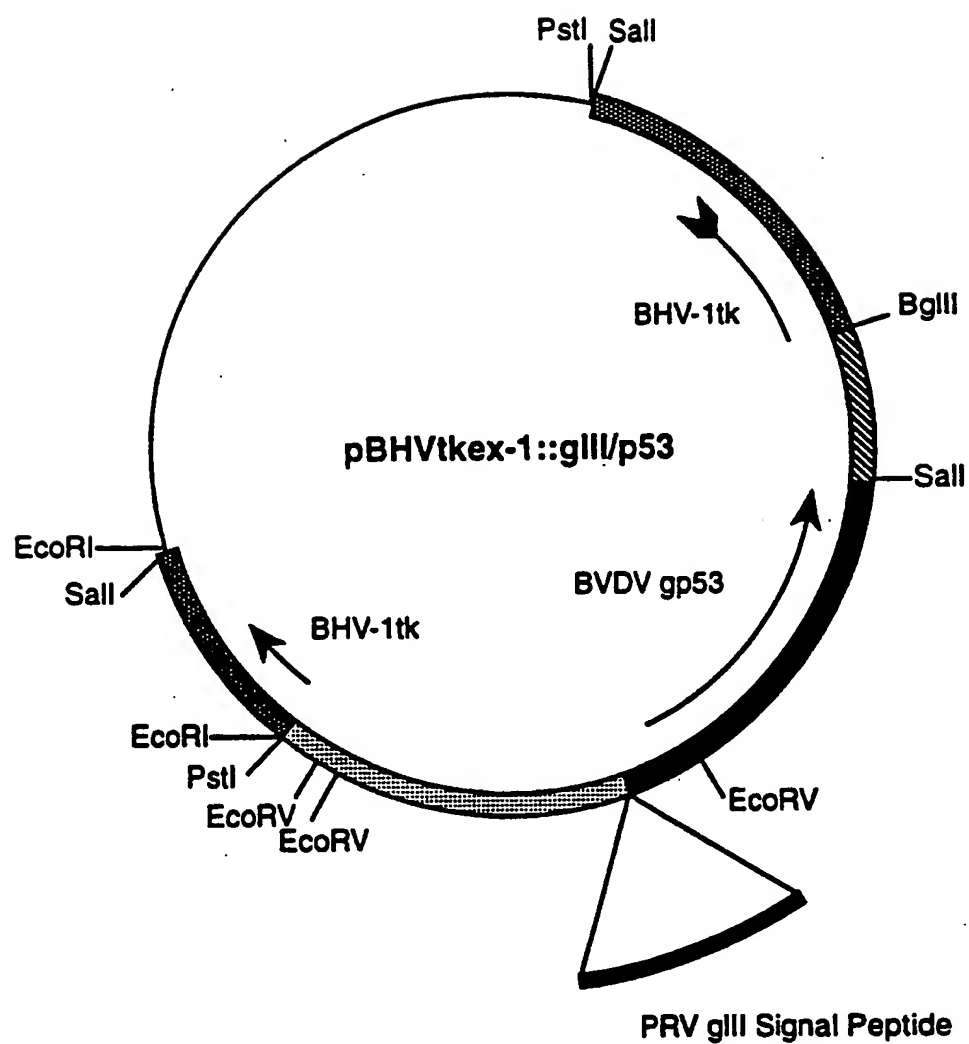
FIGURE 3C

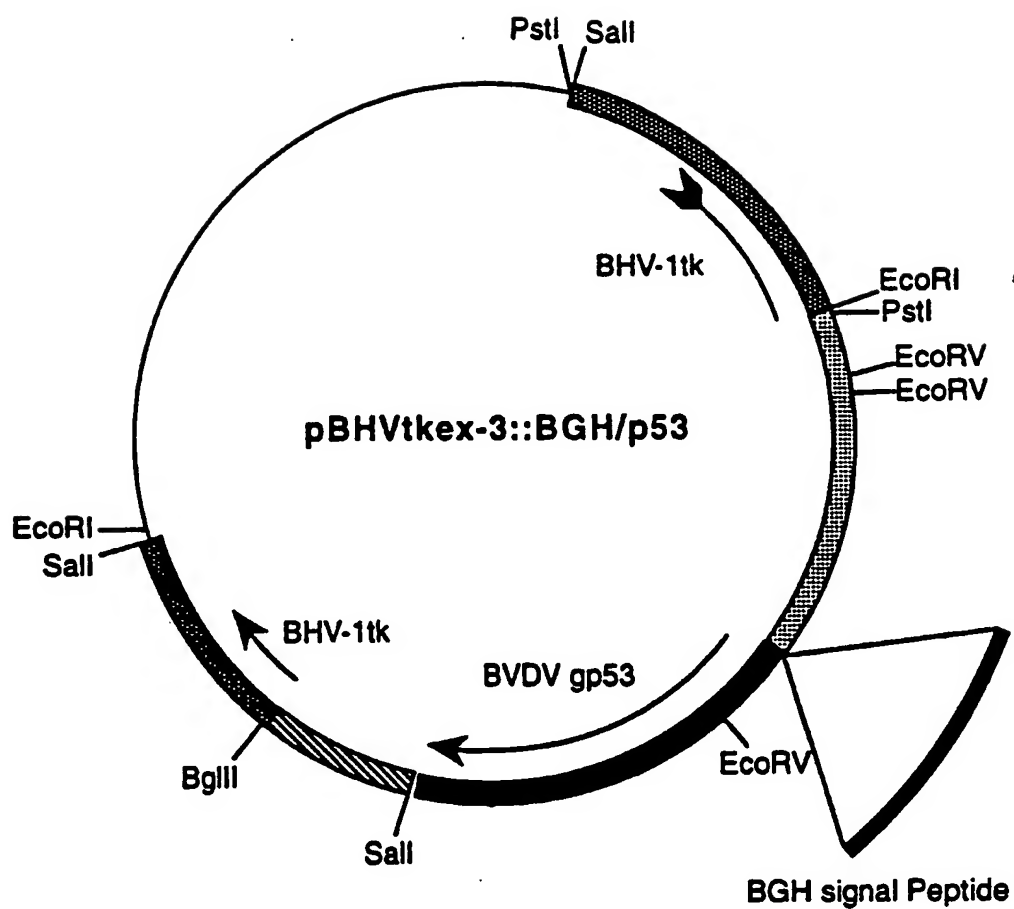
FIGURE 3D

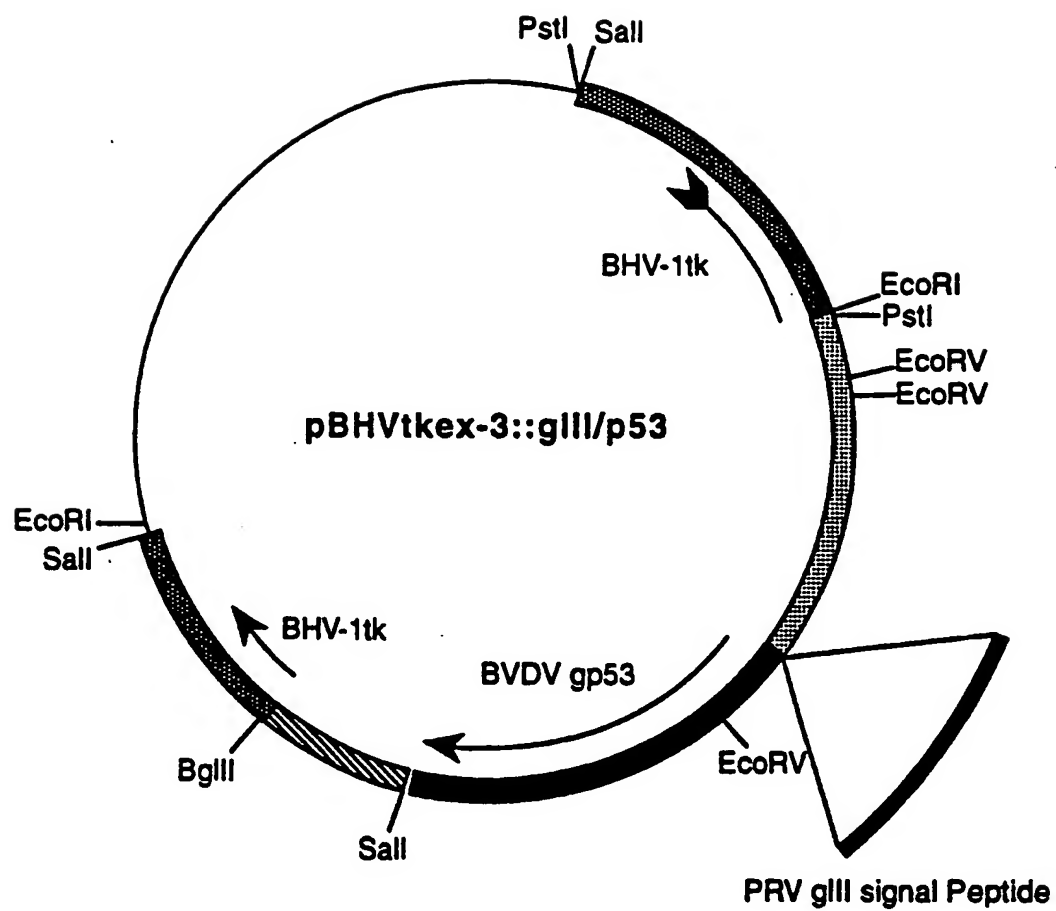
FIGURE 3E

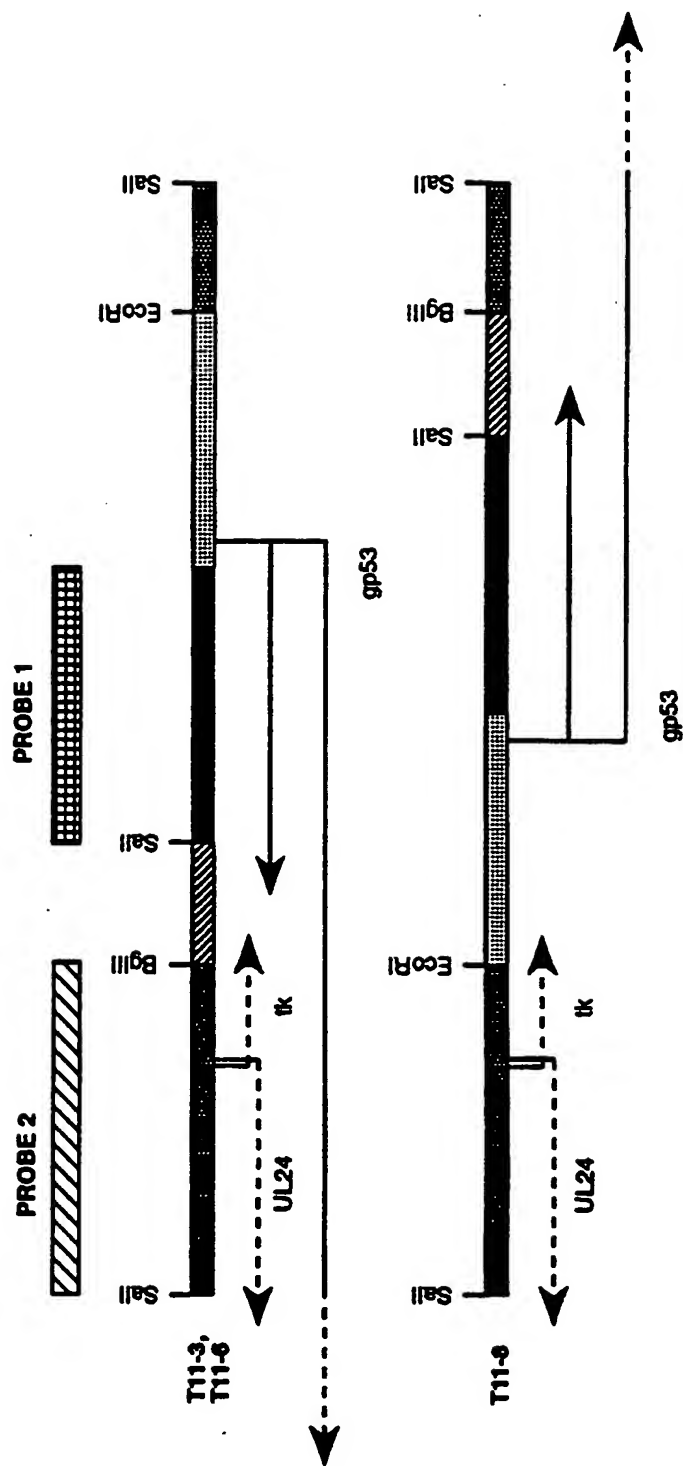
FIGURE 4

FIGURE 5A

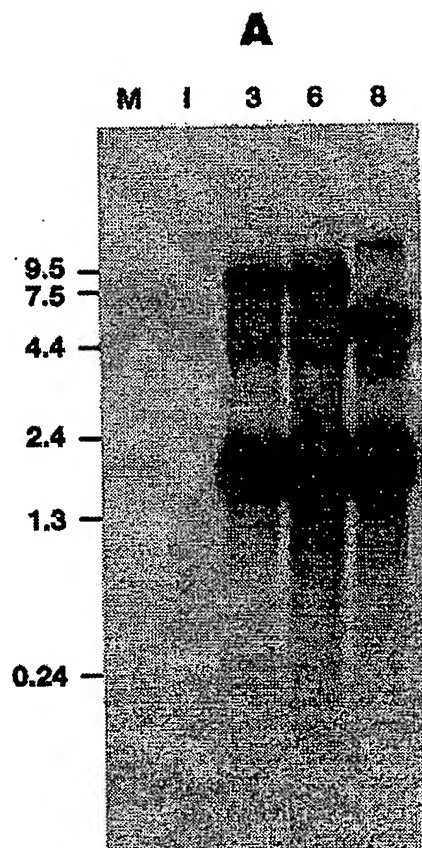


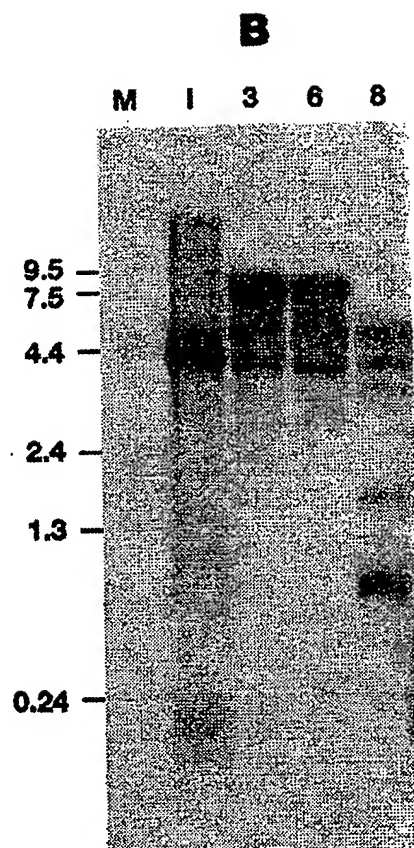
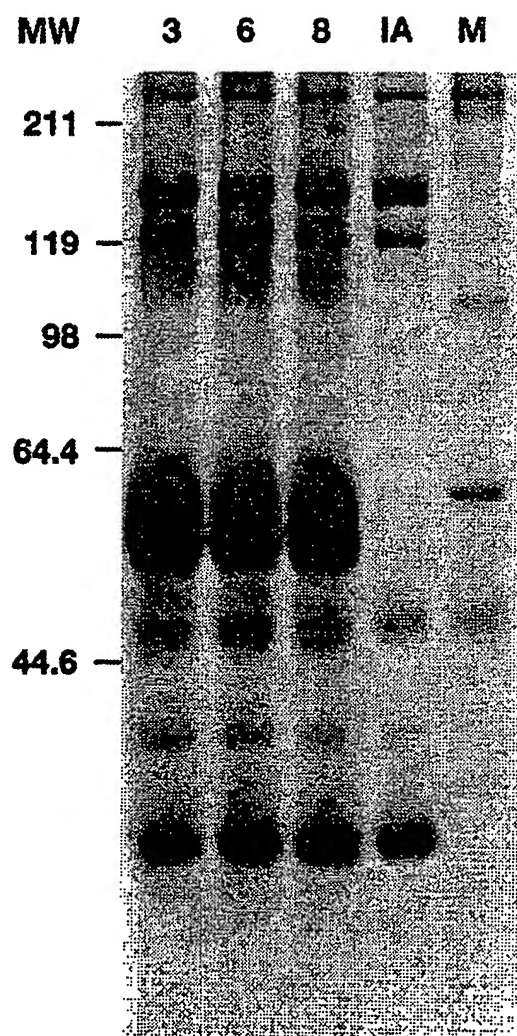
FIGURE 5B

FIGURE 6



INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification ⁶ : C12N 15/86, 7/01, C07K 14/47, 14/18, A61K 39/12	A3	(11) International Publication Number: WO 95/12682 (43) International Publication Date: 11 May 1995 (11.05.95)
(21) International Application Number: PCT/US94/12198 (22) International Filing Date: 31 October 1994 (31.10.94) (30) Priority Data: 08/147,810 5 November 1993 (05.11.93) US (60) Parent Application or Grant (63) Related by Continuation US 08/147,810 (CIP) Filed on 5 November 1993 (05.11.93) (71) Applicant (for all designated States except US): THE UPJOHN COMPANY [US/US]; 301 Henrietta Street, Kalamazoo, MI 49001 (US). (72) Inventors; and (75) Inventors/Applicants (for US only): HAANES, Elizabeth, J. [US/US]; 2030 Paddington Road, Kalamazoo, MI 49001 (US). WARDLEY, Richard, C. [US/US]; 15216 Marshfield Road, Hickory Corners, MI 49060 (US). (74) Agent: WOOTTON, Thomas, A.; The Upjohn Company, Corporate Intellectual Property Law, 301 Henrietta Street, Kalamazoo, MI 49001 (US).		(81) Designated States: AM, AT, AU, BB, BG, BR, BY, CA, CH, CN, CZ, DE, DK, EE, ES, FI, GB, GE, HU, JP, KE, KG, KP, KR, KZ, LK, LR, LT, LU, LV, MD, MG, MN, MW, NL, NO, NZ, PL, PT, RO, RU, SD, SE, SI, SK, TJ, TT, UA, US, UZ, VN, European patent (AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG), ARIPO patent (KE, MW, SD, SZ). Published <i>With international search report.</i> <i>Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i> (88) Date of publication of the international search report: 6 July 1995 (06.07.95)
(54) Title: VIRAL VECTOR WITH BOVINE VIRAL DIARRHEA VIRUS (BVDV) ANTIGENS		
(57) Abstract <p>This invention relates to the field of Bovine Viral Diarrhea Virus (BVDV), and vaccines for the treatment thereof. This invention describes the preparation of live, attenuated Bovine Herpesvirus type 1 (BHV-1) as a virus, vaccine and vector for expression of BVDV antigens. A BVDV cDNA clone containing sequences corresponding to glycoprotein gp53 is inserted into an inactivated BHV-1 virus.</p>		

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INTERNATIONAL SEARCH REPORT

International Application No
PCT/L 94/12198

A. CLASSIFICATION OF SUBJECT MATTER
IPC 6 C12N15/86 C07K14/18 C12N7/01 A61K39/12 C07K14/47

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
IPC 6 C12N C07K

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Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P,X	WO,A,94 00586 (RHONE-MERIEUX) 6 January 1994 see claim 11 ---	1-3,25, 27,28
Y	VIROLOGY, vol.190, no.2, 1992 pages 666 - 673 BELLO, L. ET AL. 'Bovine herpesvirus 1 as a live virus vector for expression of foreign genes' see the whole document --- -/--	1-3,6,7, 25,27-30

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

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Date of the actual completion of the international search

19 May 1995

Date of mailing of the international search report

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INTERNATIONAL SEARCH REPORT

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C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	VIROLOGY, vol.190, no.2, 1992 pages 763 - 772 PATON, D.J. ET AL. 'Epitope mapping of the gp53 envelope protein of bovine viral diarrhea virus' see the whole document ---	1-3,6,7, 25,27-30
P,A	VIRUS RESEARCH, vol.34, no.2, 1994 pages 178 - 186 YU, M. ET AL. 'High level expression of the envelope glycoprotein (GP53) of bovine viral diarrhoea virus (singer) and its potential use as diagnostic reagent' see the whole document ---	1
A	EP,A,0 464 010 (STATENS VETERINÄRMEDICINSKA ANSTALT) 2 January 1992 see the whole document ---	1
A	WO,A,90 01337 (INSTITUTE FOR ANIMAL HEALTH LIMITED) 22 February 1990 see the whole document ---	1
A	EP,A,0 119 025 (BAYLOR COLLEGE OF MEDECINE. NOVAGENE LTD) 19 September 1984 see the whole document -----	1

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 94/ 12198

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:
Remark: Although claim 27 is directed to a method of treatment of the animal body the search has been carried out and based on the alleged effects of the compound/composition.
2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/CC 94/12198

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO-A-9400586	06-01-94	FR-A- 2693472 AU-B- 4334193 CA-A- 2116355 EP-A- 0605680	14-01-94 24-01-94 06-01-94 13-07-94
EP-A-0464010	02-01-92	NONE	
WO-A-9001337	22-02-90	AU-B- 628845 AU-A- 4049189 CA-A- 1319634 EP-A- 0427767 GB-A- 2239799 JP-T- 4500069	24-09-92 05-03-90 29-06-93 22-05-91 17-07-91 09-01-92
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